

SEQUENCE LISTING

<110> Carozzi, Nadine
Hargiss, Tracy
Koziel, Michael G.
Duck, Nicholas B.
Carr, Brian

<120> AXMI-014, A Delta-Endotoxin Gene and
Methods for Its Use

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<151> 2003-02-20

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<222> (1)...(2019)

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aag gat cca aat ata ttt cct att aac ctg gac gct tgt cag gga agg 144
Lys Asp Pro Asn Ile Phe Pro Ile Asn Leu Asp Ala Cys Gln Gly Arg
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cca tgg caa gat acg tgg gaa tca gtc tcg gat ata gta act att ggg 192
Pro Trp Gln Asp Thr Trp Glu Ser Val Ser Asp Ile Val Thr Ile Gly
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aca tac ctt ata caa ttc ttg cta gaa ccc ggt ata ggt gga att cct 240
Thr Tyr Leu Ile Gln Phe Leu Leu Glu Pro Gly Ile Gly Gly Ile Pro
65 70 75 80

gta ata ttt tca ata ata aac aaa ctc att ccg tct tct ggt caa tct 288
Val Ile Phe Ser Ile Ile Asn Lys Leu Ile Pro Ser Ser Gly Gln Ser
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Val Ala Ala Leu Ser Ile Cys Asp Leu Val Ser Ile Ile Arg Lys Glu
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Val Asp Glu Ser Val Leu Ser Asp Gly Val Ala Asp Phe Glu Gly Glu
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Asn	Val	His	Leu	Leu	Leu	Leu	Arg	Asp	Ala	Val	Lys	Tyr	Lys	Lys	Glu	
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Trp	Gly	Leu	Val	Cys	Pro	Pro	Leu	Tyr	Pro	Gly	Ser	Gly	Arg	Thr	Asp	
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Arg His Leu Val Ser Arg Ala Glu Phe Phe Pro Thr Thr Leu Asn Thr			
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Phe Leu Tyr Glu Val Asn Ser Ser Gly Tyr Ser Gln Thr Ile Glu Ser			
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Gly Tyr Ser Ser Phe Asp Tyr Val Asp Thr Leu Val Thr Thr Phe Asn			
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Pro Asn Asn Thr Asn Met Ser Asn Arg Tyr Pro Phe Ala Lys Asp Pro
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aat ata ttt cct att aac ctg gac gct tgt cag gga agg cca tgg caa 144
Asn Ile Phe Pro Ile Asn Leu Asp Ala Cys Gln Gly Arg Pro Trp Gln
35 40 45

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Asp Thr Trp Glu Ser Val Ser Asp Ile Val Thr Ile Gly Thr Tyr Leu
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Ile Gln Phe Leu Leu Glu Pro Gly Ile Gly Gly Ile Pro Val Ile Phe
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85 90 95

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Leu Ser Ile Cys Asp Leu Val Ser Ile Ile Arg Lys Glu Val Asp Glu
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Ser Val Leu Ser Asp Gly Val Ala Asp Phe Glu Gly Glu Met Thr Ala
115 120 125

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Tyr Gln Asp Tyr Tyr Leu His Tyr Leu Glu Asp Trp Leu Thr Asp Lys
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Glu Glu Asp Phe Thr Lys Leu Leu Ala Gly Ser Leu Ser Arg Gln Lys
165 170 175

gct gaa ata tta tta ttg cct acg tat gtg caa gct gca aat gtg cat 576
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180 185 190

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Glu Val Asn Ser Ser Gly Tyr Ser Gln Thr Ile Glu Ser Val Leu Pro	
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Gly Ile Asn Lys Asp Leu Pro Pro Ser Arg Thr Asn Tyr Ser His Arg	
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Gly Thr Gly Tyr Ala Gly Gly Tyr Val Thr Ala Gly Pro Gly Tyr Thr	
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Lys Val Glu Phe Ile Pro Ile Asp Ile Gln Ile Glu Lys Cys Thr Lys	
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Cys Gln Phe Glu Gly Asp Ile Cys Arg Cys Glu Gly Val Gln Ser Leu
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Ala	Glu	Lys	Lys	Trp	Arg	Asp	Lys	Arg	Glu	Lys	Leu	Glu	Trp	Glu	Thr
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Arg	Gly	Tyr	Ile	Glu	Asp	Ser	Gln	Asp	Leu	Glu	Ile	Tyr	Leu	Ile
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Trp	Pro	Leu	Ser	Ala	Gln	Ser	Pro	Ile	Gly	Lys	Cys	Gly	Glu	Pro
785					790					795				800
Arg	Cys	Ala	Pro	His	Leu	Glu	Trp	Asn	Pro	Asp	Leu	Asp	Cys	Ser
				805				810					815	
Arg	Asp	Gly	Glu	Lys	Cys	Ala	His	His	Ser	His	His	Phe	Ser	Leu
			820					825					830	
Ile	Asp	Val	Gly	Cys	Thr	Asp	Leu	Asn	Glu	Asp	Leu	Gly	Val	Trp
		835					840					845		
Ile	Phe	Lys	Ile	Lys	Thr	Gln	Asp	Gly	His	Ala	Arg	Leu	Gly	Asn
	850					855					860			
Glu	Phe	Leu	Glu	Glu	Lys	Pro	Leu	Val	Gly	Glu	Ala	Leu	Ala	Arg
865					870					875				880
Lys	Arg	Ala	Glu	Lys	Trp	Arg	Asp	Lys	Arg	Glu	Lys	Leu	Glu	Trp
				885				890						895
Glu	Thr	Asn	Ile	Val	Tyr	Lys	Glu	Ala	Lys	Glu	Ser	Val	Asp	Ala
		900						905					910	
Phe	Val	Asn	Ser	Gln	Tyr	Asp	Gln	Leu	Gln	Ala	Asp	Thr	Asn	Ile
		915					920					925		
Met	Ile	His	Ala	Ala	Asp	Lys	Arg	Val	His	Ser	Ile	Arg	Glu	Ala
	930					935					940			
Leu	Pro	Glu	Leu	Ser	Val	Ile	Pro	Gly	Val	Asn	Ala	Ala	Ile	Phe
945					950					955				960
Glu	Leu	Glu	Gly	Arg	Ile	Phe	Thr	Ala	Phe	Ser	Leu	Tyr	Asp	Ala
				965					970					975
Asn	Val	Ile	Lys	Asn	Gly	Asp	Phe	Asn	Asn	Gly	Leu	Ser	Cys	Trp
			980				985						990	
Val	Lys	Gly	His	Val	Asp	Val	Glu	Glu	Gln	Asn	Asn	Gln	Arg	Ser
		995					1000					1005		
Leu	Val	Val	Pro	Glu	Trp	Glu	Ala	Glu	Val	Ser	Gln	Glu	Val	Arg
	1010					1015						1020		
Cys	Pro	Gly	Arg	Gly	Tyr	Ile	Leu	Arg	Val	Thr	Ala	Tyr	Lys	Glu
1025					1030					1035				1040
Tyr	Gly	Glu	Gly	Cys	Val	Thr	Ile	His	Glu	Ile	Glu	Asn	Asn	Thr
				1045					1050					1055
Glu	Leu	Lys	Phe	Ser	Asn	Cys	Val	Glu	Glu	Glu	Ile	Tyr	Pro	Asn
			1060					1065					1070	
Thr	Val	Thr	Cys	Asn	Asp	Tyr	Thr	Val	Asn	Gln	Glu	Glu	Tyr	Gly
		1075					1080						1085	

Ala	Tyr	Thr	Ser	Arg	Asn	Arg	Gly	Tyr	Asn	Glu	Ala	Pro	Ser	Val	Pro	
1090						1095				1100						
Ala	Asp	Tyr	Ala	Ser	Val	Tyr	Glu	Glu	Lys	Ser	Tyr	Thr	Asp	Gly	Arg	
1105					1110					1115					1120	
Arg	Glu	Asn	Pro	Cys	Glu	Phe	Asn	Arg	Gly	Tyr	Arg	Asp	Tyr	Thr	Pro	
				1125					1130						1135	
Leu	Pro	Val	Gly	Tyr	Val	Thr	Lys	Glu	Leu	Glu	Tyr	Phe	Pro	Glu	Thr	
			1140					1145					1150			
Asp	Lys	Val	Trp	Ile	Glu	Ile	Gly	Glu	Thr	Glu	Gly	Thr	Phe	Ile	Val	
		1155					1160					1165				
Asp	Ser	Val	Glu	Leu	Leu	Leu	Met	Glu	Glu							
	1170						1175									

<210> 8
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 <212> PRT
 <213> Bacillus thuringiensis

<400> 8																
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Lys	Val	Asp	Lys	Ile	Ser	Thr	Asp	Ser	Leu	Lys	Asn	Glu	Thr	Asp	Ile	
			20					25					30			
Glu	Leu	Gln	Asn	Ile	Asn	His	Glu	Asp	Cys	Leu	Lys	Met	Ser	Glu	Tyr	
		35					40					45				
Glu	Asn	Val	Glu	Pro	Phe	Val	Ser	Ala	Ser	Thr	Ile	Gln	Thr	Gly	Ile	
	50					55					60					
Gly	Ile	Ala	Gly	Lys	Ile	Leu	Gly	Thr	Leu	Gly	Val	Pro	Phe	Ala	Gly	
65				70					75						80	
Gln	Val	Ala	Ser	Leu	Tyr	Ser	Phe	Ile	Leu	Gly	Glu	Leu	Trp	Pro	Lys	
				85					90					95		
Gly	Lys	Asn	Gln	Trp	Glu	Ile	Phe	Met	Glu	His	Val	Glu	Glu	Ile	Ile	
			100					105					110			
Asn	Gln	Lys	Ile	Ser	Thr	Tyr	Ala	Arg	Asn	Lys	Ala	Leu	Thr	Asp	Leu	
		115					120					125				
Lys	Gly	Leu	Gly	Asp	Ala	Leu	Ala	Val	Tyr	His	Asp	Ser	Leu	Glu	Ser	
	130					135					140					
Trp	Val	Gly	Asn	Arg	Asn	Asn	Thr	Arg	Ala	Arg	Ser	Val	Val	Lys	Ser	
145				150					155						160	
Gln	Tyr	Ile	Ala	Leu	Glu	Leu	Met	Phe	Val	Gln	Lys	Leu	Pro	Ser	Phe	
				165				170						175		
Ala	Val	Ser	Gly	Glu	Glu	Val	Pro	Leu	Leu	Pro	Ile	Tyr	Ala	Gln	Ala	
			180					185					190			
Ala	Asn	Leu	His	Leu	Leu	Leu	Leu	Arg	Asp	Ala	Ser	Ile	Phe	Gly	Lys	
	195						200					205				
Glu	Trp	Gly	Leu	Ser	Ser	Ser	Glu	Ile	Ser	Thr	Phe	Tyr	Asn	Arg	Gln	
	210					215					220					
Val	Glu	Arg	Ala	Gly	Asp	Tyr	Ser	Asp	His	Cys	Val	Lys	Trp	Tyr	Ser	
225				230					235						240	
Thr	Gly	Leu	Asn	Asn	Leu	Arg	Gly	Thr	Asn	Ala	Glu	Ser	Trp	Val	Arg	
				245				250						255		
Tyr	Asn	Gln	Phe	Arg	Arg	Asp	Met	Thr	Leu	Met	Val	Leu	Asp	Leu	Val	
		260					265						270			
Ala	Leu	Phe	Pro	Ser	Tyr	Asp	Thr	Gln	Met	Tyr	Pro	Ile	Lys	Thr	Thr	
	275						280					285				
Ala	Gln	Leu	Thr	Arg	Glu	Val	Tyr	Thr	Asp	Ala	Ile	Gly	Thr	Val	His	

290		295		300
Pro His Pro Ser Phe Thr Ser Thr Thr Trp Tyr Asn Asn Asn Ala Pro				
305		310		315
Ser Phe Ser Ala Ile Glu Ala Ala Val Val Arg Asn Pro His Leu Leu				
		325		330
Asp Phe Leu Glu Gln Val Thr Ile Tyr Ser Leu Leu Ser Arg Trp Ser				
		340		345
Asn Thr Gln Tyr Met Asn Met Trp Gly Gly His Lys Leu Glu Phe Arg				
		355		360
Thr Ile Gly Gly Thr Leu Asn Ile Ser Thr Gln Gly Ser Thr Asn Thr				
		370		375
Ser Ile Asn Pro Val Thr Leu Pro Phe Thr Ser Arg Asp Val Tyr Arg				
385		390		395
Thr Glu Ser Leu Ala Gly Leu Asn Leu Phe Leu Thr Gln Pro Val Asn				
		405		410
Gly Val Pro Arg Val Asp Phe His Trp Lys Phe Val Thr His Pro Ile				
		420		425
Ala Ser Asp Asn Phe Tyr Tyr Pro Gly Tyr Ala Gly Ile Gly Thr Gln				
		435		440
Leu Gln Asp Ser Glu Asn Glu Leu Pro Pro Glu Ala Thr Gly Gln Pro				
		450		455
Asn Tyr Glu Ser Tyr Ser His Arg Leu Ser His Ile Gly Leu Ile Ser				
465		470		475
Ala Ser His Val Lys Ala Leu Val Tyr Ser Trp Thr His Arg Ser Ala				
		485		490
Asp Arg Thr Asn Thr Ile Glu Pro Asn Ser Ile Thr Gln Ile Pro Leu				
		500		505
Val Lys Ala Phe Asn Leu Ser Ser Gly Ala Ala Val Val Arg Gly Pro				
		515		520
Gly Phe Thr Gly Gly Asp Ile Leu Arg Arg Thr Asn Thr Gly Thr Phe				
		530		535
Gly Asp Ile Arg Val Asn Ile Asn Pro Pro Phe Ala Gln Arg Tyr Arg				
545		550		555
Val Arg Ile Arg Tyr Ala Ser Thr Thr Asp Leu Gln Phe His Thr Ser				
		565		570
Ile Asn Gly Lys Ala Ile Asn Gln Gly Asn Phe Ser Ala Thr Met Asn				
		580		585
Arg Gly Glu Asp Leu Asp Tyr Lys Thr Phe Arg Thr Val Gly Phe Thr				
		595		600
Thr Pro Phe Ser Phe Leu Asp Val Gln Ser Thr Phe Thr Ile Gly Ala				
		610		615
Trp Asn Phe Ser Ser Gly Asn Glu Val Tyr Ile Asp Arg Ile Glu Phe				
625		630		635
Val Pro Val Glu Val Thr Tyr Glu Ala Glu Tyr Asp Phe Glu Lys Ala				
		645		650
Gln Glu Lys Val Thr Ala Leu Phe Thr Ser Thr Asn Pro Arg Gly Leu				
		660		665
Lys Thr Asp Val Lys Asp Tyr His Ile Asp Gln Val Ser Asn Leu Val				
		675		680
Glu Ser Leu Ser Asp Glu Phe Tyr Leu Asp Glu Lys Arg Glu Leu Phe				
		690		695
Glu Ile Val Lys Tyr Ala Lys Gln Leu His Ile Glu Arg Asn Met				
705		710		715

<210> 9
<211> 633

<212> PRT

<213> *Bacillus thuringiensis*

<400> 9

Met	Asn	Asn	Val	Leu	Asn	Ser	Gly	Arg	Thr	Thr	Ile	Cys	Asp	Ala	Tyr
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Asn	Val	Val	Ala	His	Asp	Pro	Phe	Ser	Phe	Glu	His	Lys	Ser	Leu	Asp
			20					25					30		
Thr	Ile	Gln	Lys	Glu	Trp	Met	Glu	Trp	Lys	Arg	Thr	Asp	His	Ser	Leu
		35					40					45			
Tyr	Val	Ala	Pro	Val	Val	Gly	Thr	Val	Ser	Ser	Phe	Leu	Leu	Lys	Lys
	50					55					60				
Val	Gly	Ser	Leu	Ile	Gly	Lys	Arg	Ile	Leu	Ser	Glu	Leu	Trp	Gly	Ile
65					70					75					80
Ile	Phe	Pro	Ser	Gly	Ser	Thr	Asn	Leu	Met	Gln	Asp	Ile	Leu	Arg	Glu
				85					90					95	
Thr	Glu	Gln	Phe	Leu	Asn	Gln	Arg	Leu	Asn	Thr	Asp	Thr	Leu	Ala	Arg
			100					105					110		
Val	Asn	Ala	Glu	Leu	Ile	Gly	Leu	Gln	Ala	Asn	Ile	Arg	Glu	Phe	Asn
		115					120					125			
Gln	Gln	Val	Asp	Asn	Phe	Leu	Asn	Pro	Thr	Gln	Asn	Pro	Val	Pro	Leu
	130					135					140				
Ser	Ile	Thr	Ser	Ser	Val	Asn	Thr	Met	Gln	Gln	Leu	Phe	Leu	Asn	Arg
145					150					155					160
Leu	Pro	Gln	Phe	Gln	Ile	Gln	Gly	Tyr	Gln	Leu	Leu	Leu	Leu	Pro	Leu
				165					170					175	
Phe	Ala	Gln	Ala	Ala	Asn	Met	His	Leu	Ser	Phe	Ile	Arg	Asp	Val	Ile
			180					185					190		
Leu	Asn	Ala	Asp	Glu	Trp	Gly	Ile	Ser	Ala	Ala	Thr	Leu	Arg	Thr	Tyr
	195					200						205			
Arg	Asp	Tyr	Leu	Arg	Asn	Tyr	Thr	Arg	Asp	Tyr	Ser	Asn	Tyr	Cys	Ile
	210				215						220				
Asn	Thr	Tyr	Gln	Thr	Ala	Phe	Arg	Gly	Leu	Asn	Thr	Arg	Leu	His	Asp
225					230					235					240
Met	Leu	Glu	Phe	Arg	Thr	Tyr	Met	Phe	Leu	Asn	Val	Phe	Glu	Tyr	Val
			245					250					255		
Ser	Ile	Trp	Ser	Leu	Phe	Lys	Tyr	Gln	Ser	Leu	Met	Val	Ser	Ser	Gly
		260						265				270			
Ala	Asn	Leu	Tyr	Ala	Ser	Gly	Ser	Gly	Pro	Gln	Gln	Thr	Gln	Ser	Phe
		275				280						285			
Thr	Ala	Gln	Asn	Trp	Pro	Phe	Leu	Tyr	Ser	Leu	Phe	Gln	Val	Asn	Ser
	290					295					300				
Asn	Tyr	Ile	Leu	Ser	Gly	Ile	Ser	Gly	Thr	Arg	Leu	Ser	Ile	Thr	Phe
305					310					315					320
Pro	Asn	Ile	Gly	Gly	Leu	Pro	Gly	Ser	Thr	Thr	Thr	His	Ser	Leu	Asn
			325					330					335		
Ser	Ala	Arg	Val	Asn	Tyr	Ser	Gly	Gly	Val	Ser	Ser	Gly	Leu	Ile	Gly
		340						345				350			
Ala	Thr	Asn	Leu	Asn	His	Asn	Phe	Asn	Cys	Ser	Thr	Val	Leu	Pro	Pro
		355					360					365			
Leu	Ser	Thr	Pro	Phe	Val	Arg	Ser	Trp	Leu	Asp	Ser	Gly	Thr	Asp	Arg
	370					375					380				
Glu	Gly	Val	Ala	Thr	Ser	Thr	Asn	Trp	Gln	Thr	Glu	Ser	Phe	Gln	Thr
385					390					395					400
Thr	Leu	Ser	Leu	Arg	Cys	Gly	Ala	Phe	Ser	Ala	Arg	Gly	Asn	Ser	Asn
			405					410					415		
Tyr	Phe	Pro	Asp	Tyr	Phe	Ile	Arg	Asn	Ile	Ser	Gly	Val	Pro	Leu	Val

			420					425					430				
Ile	Arg	Asn	Glu	Asp	Leu	Thr	Arg	Pro	Leu	His	Tyr	Asn	Gln	Ile	Arg		
		435						440				445					
Asn	Ile	Glu	Ser	Pro	Ser	Gly	Thr	Pro	Gly	Gly	Ala	Arg	Ala	Tyr	Leu		
		450						455				460					
Val	Ser	Val	His	Asn	Arg	Lys	Asn	Asn	Ile	Tyr	Ala	Ala	Asn	Glu	Asn		
465					470					475				480			
Gly	Thr	Met	Ile	His	Leu	Ala	Pro	Glu	Asp	Tyr	Thr	Gly	Phe	Thr	Ile		
				485						490				495			
Ser	Pro	Ile	His	Ala	Thr	Gln	Val	Asn	Asn	Gln	Thr	Arg	Thr	Phe	Ile		
			500					505					510				
Ser	Glu	Lys	Phe	Gly	Asn	Gln	Gly	Asp	Ser	Leu	Arg	Phe	Glu	Gln	Ser		
		515						520				525					
Asn	Thr	Thr	Ala	Arg	Tyr	Thr	Leu	Arg	Gly	Asn	Gly	Asn	Ser	Tyr	Asn		
		530					535				540						
Leu	Tyr	Leu	Arg	Val	Ser	Ser	Ile	Gly	Asn	Ser	Thr	Ile	Arg	Val	Thr		
545					550					555				560			
Ile	Asn	Gly	Arg	Val	Tyr	Thr	Val	Ser	Asn	Val	Asn	Thr	Thr	Thr	Asn		
				565					570					575			
Asn	Asp	Gly	Val	Asn	Asp	Asn	Gly	Ala	Arg	Phe	Ser	Asp	Ile	Asn	Ile		
			580					585				590					
Gly	Asn	Ile	Val	Ala	Ser	Asp	Asn	Thr	Asn	Val	Thr	Leu	Asp	Ile	Asn		
		595					600					605					
Val	Thr	Leu	Asn	Ser	Gly	Thr	Pro	Phe	Asp	Leu	Met	Asn	Ile	Met	Phe		
		610					615				620						
Val	Pro	Thr	Asn	Leu	Pro	Pro	Leu	Tyr									
625							630										

<210> 10

<211> 652

<212> PRT

<213> Bacillus thuringiensis

<400> 10

Met	Ile	Arg	Lys	Gly	Gly	Arg	Lys	Met	Asn	Pro	Asn	Asn	Arg	Ser	Glu		
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His	Asp	Thr	Ile	Lys	Thr	Thr	Glu	Asn	Asn	Glu	Val	Pro	Thr	Asn	His		
			20					25					30				
Val	Gln	Tyr	Pro	Leu	Ala	Glu	Thr	Pro	Asn	Pro	Thr	Leu	Glu	Asp	Leu		
		35					40					45					
Asn	Tyr	Lys	Glu	Phe	Leu	Arg	Met	Thr	Ala	Asp	Asn	Asn	Thr	Glu	Ala		
		50				55				60							
Leu	Asp	Ser	Ser	Thr	Thr	Lys	Asp	Val	Ile	Gln	Lys	Gly	Ile	Ser	Val		
65					70					75				80			
Val	Gly	Asp	Leu	Leu	Gly	Val	Val	Gly	Phe	Pro	Phe	Gly	Gly	Ala	Leu		
			85					90					95				
Val	Ser	Phe	Tyr	Thr	Asn	Phe	Leu	Asn	Thr	Ile	Trp	Pro	Ser	Glu	Asp		
		100						105					110				
Pro	Trp	Lys	Ala	Phe	Met	Glu	Gln	Val	Glu	Ala	Leu	Met	Asp	Gln	Lys		
		115					120					125					
Ile	Ala	Asp	Tyr	Ala	Lys	Asn	Lys	Ala	Leu	Ala	Glu	Leu	Gln	Gly	Leu		
		130				135					140						
Gln	Asn	Asn	Val	Glu	Asp	Tyr	Val	Ser	Ala	Leu	Ser	Ser	Trp	Gln	Lys		
145					150					155				160			
Asn	Pro	Val	Ser	Ser	Arg	Asn	Pro	His	Ser	Gln	Gly	Arg	Ile	Arg	Glu		
				165					170					175			

Leu	Phe	Ser	Gln	Ala	Glu	Ser	His	Phe	Arg	Asn	Ser	Met	Pro	Ser	Phe
			180					185					190		
Ala	Ile	Ser	Gly	Tyr	Glu	Val	Leu	Phe	Leu	Thr	Thr	Tyr	Ala	Gln	Ala
		195					200					205			
Ala	Asn	Thr	His	Leu	Phe	Leu	Leu	Lys	Asp	Ala	Gln	Ile	Tyr	Gly	Glu
	210					215					220				
Glu	Trp	Gly	Tyr	Glu	Lys	Glu	Asp	Ile	Ala	Glu	Phe	Tyr	Lys	Arg	Gln
225					230					235					240
Leu	Lys	Leu	Thr	Gln	Glu	Tyr	Thr	Asp	His	Cys	Val	Lys	Trp	Tyr	Asn
			245					250						255	
Val	Gly	Leu	Asp	Lys	Leu	Arg	Gly	Ser	Ser	Tyr	Glu	Ser	Trp	Val	Asn
		260						265					270		
Phe	Asn	Arg	Tyr	Arg	Arg	Glu	Met	Thr	Leu	Thr	Val	Leu	Asp	Leu	Ile
	275						280					285			
Ala	Leu	Phe	Pro	Leu	Tyr	Asp	Val	Arg	Leu	Tyr	Pro	Lys	Glu	Val	Lys
	290					295					300				
Thr	Glu	Leu	Thr	Arg	Asp	Val	Leu	Thr	Asp	Pro	Ile	Val	Gly	Val	Asn
305					310					315					320
Asn	Leu	Arg	Gly	Tyr	Gly	Thr	Thr	Phe	Ser	Asn	Ile	Glu	Asn	Tyr	Ile
			325					330						335	
Arg	Lys	Pro	His	Leu	Phe	Asp	Tyr	Leu	His	Arg	Ile	Gln	Phe	His	Thr
		340						345					350		
Arg	Phe	Gln	Pro	Gly	Tyr	Tyr	Gly	Asn	Asp	Ser	Phe	Asn	Tyr	Trp	Ser
	355						360					365			
Gly	Asn	Tyr	Val	Ser	Thr	Arg	Pro	Ser	Ile	Gly	Ser	Asn	Asp	Ile	Ile
	370					375					380				
Thr	Ser	Pro	Phe	Tyr	Gly	Asn	Lys	Ser	Ser	Glu	Pro	Val	Gln	Asn	Leu
385					390					395					400
Glu	Phe	Asn	Gly	Glu	Lys	Val	Tyr	Arg	Ala	Val	Ala	Asn	Thr	Asn	Leu
			405						410					415	
Ala	Val	Trp	Pro	Ser	Ala	Val	Tyr	Ser	Gly	Val	Thr	Lys	Val	Glu	Phe
		420						425					430		
Ser	Gln	Tyr	Asn	Asp	Gln	Thr	Asp	Glu	Ala	Ser	Thr	Gln	Thr	Tyr	Asp
	435						440					445			
Ser	Lys	Arg	Asn	Val	Gly	Ala	Val	Ser	Trp	Asp	Ser	Ile	Asp	Gln	Leu
	450					455					460				
Pro	Pro	Glu	Thr	Thr	Asp	Glu	Pro	Leu	Glu	Lys	Gly	Tyr	Ser	His	Gln
465					470					475					480
Leu	Asn	Tyr	Val	Met	Cys	Phe	Leu	Met	Gln	Gly	Ser	Arg	Gly	Thr	Ile
			485						490					495	
Pro	Val	Leu	Thr	Trp	Thr	His	Lys	Ser	Val	Asp	Phe	Phe	Asn	Met	Ile
		500						505					510		
Asp	Ser	Lys	Lys	Ile	Thr	Gln	Leu	Pro	Leu	Val	Lys	Ala	Tyr	Lys	Leu
	515						520					525			
Gln	Ser	Gly	Ala	Ser	Val	Val	Ala	Gly	Pro	Arg	Phe	Thr	Gly	Gly	Asp
	530					535					540				
Ile	Ile	Gln	Cys	Thr	Glu	Asn	Gly	Ser	Ala	Ala	Thr	Ile	Tyr	Val	Thr
545					550					555					560
Pro	Asp	Val	Ser	Tyr	Ser	Gln	Lys	Tyr	Arg	Ala	Arg	Ile	His	Tyr	Ala
			565						570					575	
Ser	Thr	Ser	Gln	Ile	Thr	Phe	Thr	Leu	Ser	Leu	Asp	Gly	Ala	Pro	Phe
		580						585					590		
Asn	Gln	Tyr	Tyr	Phe	Asp	Lys	Thr	Ile	Asn	Lys	Gly	Asp	Thr	Leu	Thr
	595						600					605			
Tyr	Asn	Ser	Phe	Asn	Leu	Ala	Ser	Phe	Ser	Thr	Pro	Phe	Glu	Leu	Ser
	610					615					620				
Gly	Asn	Asn	Leu	Gln	Ile	Gly	Val	Thr	Gly	Leu	Ser	Ala	Gly	Asp	Lys

625		630		635	640						
Val	Tyr	Ile	Asp	Lys	Ile	Glu	Phe	Ile	Pro	Val	Asn
		645		650							

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 <212> PRT
 <213> *Bacillus thuringiensis*

<400> 11

Met	Asn	Pro	Asn	Asn	Arg	Ser	Glu	His	Asp	Thr	Ile	Lys	Val	Thr	Pro
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Asn	Ser	Glu	Leu	Gln	Thr	Asn	His	Asn	Gln	Tyr	Pro	Leu	Ala	Asp	Asn
			20					25					30		
Pro	Asn	Ser	Thr	Leu	Glu	Glu	Leu	Asn	Tyr	Lys	Glu	Phe	Leu	Arg	Met
		35					40					45			
Thr	Glu	Asp	Ser	Ser	Thr	Glu	Val	Leu	Asp	Asn	Ser	Thr	Val	Lys	Asp
	50					55					60				
Ala	Val	Gly	Thr	Gly	Ile	Ser	Val	Val	Gly	Gln	Ile	Leu	Gly	Val	Val
65				70					75					80	
Gly	Val	Pro	Phe	Ala	Gly	Ala	Leu	Thr	Ser	Phe	Tyr	Gln	Ser	Phe	Leu
			85					90					95		
Asn	Thr	Ile	Trp	Pro	Ser	Asp	Ala	Asp	Pro	Trp	Lys	Ala	Phe	Met	Ala
			100					105					110		
Gln	Val	Glu	Val	Leu	Ile	Asp	Lys	Lys	Ile	Glu	Glu	Tyr	Ala	Lys	Ser
	115					120						125			
Lys	Ala	Leu	Ala	Glu	Leu	Gln	Gly	Leu	Gln	Asn	Asn	Phe	Glu	Asp	Tyr
	130					135					140				
Val	Asn	Ala	Leu	Asn	Ser	Trp	Lys	Lys	Thr	Pro	Leu	Ser	Leu	Arg	Ser
145				150						155				160	
Lys	Arg	Ser	Gln	Asp	Arg	Ile	Arg	Glu	Leu	Phe	Ser	Gln	Ala	Glu	Ser
			165					170						175	
His	Phe	Arg	Asn	Ser	Met	Pro	Ser	Phe	Ala	Val	Ser	Lys	Phe	Glu	Val
			180					185					190		
Leu	Phe	Leu	Pro	Thr	Tyr	Ala	Gln	Ala	Ala	Asn	Thr	His	Leu	Leu	Leu
		195					200					205			
Leu	Lys	Asp	Ala	Gln	Val	Phe	Gly	Glu	Glu	Trp	Gly	Tyr	Ser	Ser	Glu
	210					215					220				
Asp	Val	Ala	Glu	Phe	Tyr	His	Arg	Gln	Leu	Lys	Leu	Thr	Gln	Gln	Tyr
225				230						235				240	
Thr	Asp	His	Cys	Val	Asn	Trp	Tyr	Asn	Val	Gly	Leu	Asn	Gly	Leu	Arg
			245					250					255		
Gly	Ser	Thr	Tyr	Asp	Ala	Trp	Val	Lys	Phe	Asn	Arg	Phe	Arg	Arg	Glu
		260						265					270		
Met	Thr	Leu	Thr	Val	Leu	Asp	Leu	Ile	Val	Leu	Phe	Pro	Phe	Tyr	Asp
		275				280						285			
Ile	Arg	Leu	Tyr	Ser	Lys	Gly	Val	Lys	Thr	Glu	Leu	Thr	Arg	Asp	Ile
	290					295					300				
Phe	Thr	Asp	Pro	Ile	Phe	Ser	Leu	Asn	Thr	Leu	Gln	Glu	Tyr	Gly	Pro
305				310						315				320	
Thr	Phe	Leu	Ser	Ile	Glu	Asn	Ser	Ile	Arg	Lys	Pro	His	Leu	Phe	Asp
			325					330					335		
Tyr	Leu	Gln	Gly	Ile	Glu	Phe	His	Thr	Arg	Leu	Gln	Pro	Gly	Tyr	Phe
		340						345					350		
Gly	Lys	Asp	Ser	Phe	Asn	Tyr	Trp	Ser	Gly	Asn	Tyr	Val	Glu	Thr	Arg
	355						360					365			

Pro	Ser	Ile	Gly	Ser	Ser	Lys	Thr	Ile	Thr	Ser	Pro	Phe	Tyr	Gly	Asp
	370					375					380				
Lys	Ser	Thr	Glu	Pro	Val	Gln	Lys	Leu	Ser	Phe	Asp	Gly	Gln	Lys	Val
385					390					395					400
Tyr	Arg	Thr	Ile	Ala	Asn	Thr	Asp	Val	Ala	Ala	Trp	Pro	Asn	Gly	Lys
				405					410					415	
Val	Tyr	Leu	Gly	Val	Thr	Lys	Val	Asp	Phe	Ser	Gln	Tyr	Asp	Asp	Gln
			420					425					430		
Lys	Asn	Glu	Thr	Ser	Thr	Gln	Thr	Tyr	Asp	Ser	Lys	Arg	Asn	Asn	Gly
		435					440					445			
His	Val	Ser	Ala	Gln	Asp	Ser	Ile	Asp	Gln	Leu	Pro	Pro	Glu	Thr	Thr
	450					455					460				
Asp	Glu	Pro	Leu	Glu	Lys	Ala	Tyr	Ser	His	Gln	Leu	Asn	Tyr	Ala	Glu
465					470					475					480
Cys	Phe	Leu	Met	Gln	Asp	Arg	Arg	Gly	Thr	Ile	Pro	Phe	Phe	Thr	Trp
				485					490					495	
Thr	His	Arg	Ser	Val	Asp	Phe	Phe	Asn	Thr	Ile	Asp	Ala	Glu	Lys	Ile
			500					505					510		
Thr	Gln	Leu	Pro	Val	Val	Lys	Ala	Tyr	Ala	Leu	Ser	Ser	Gly	Ala	Ser
		515					520					525			
Ile	Ile	Glu	Gly	Pro	Gly	Phe	Thr	Gly	Gly	Asn	Leu	Leu	Phe	Leu	Lys
	530					535					540				
Glu	Ser	Ser	Asn	Ser	Ile	Ala	Lys	Phe	Lys	Val	Thr	Leu	Asn	Ser	Ala
545					550					555					560
Ala	Leu	Leu	Gln	Arg	Tyr	Arg	Val	Arg	Ile	Arg	Tyr	Ala	Ser	Thr	Thr
				565					570					575	
Asn	Leu	Arg	Leu	Phe	Val	Gln	Asn	Ser	Asn	Asn	Asp	Phe	Leu	Val	Ile
			580					585					590		
Tyr	Ile	Asn	Lys	Thr	Met	Asn	Lys	Asp	Asp	Asp	Leu	Thr	Tyr	Gln	Thr
		595					600					605			
Phe	Asp	Leu	Ala	Thr	Thr	Asn	Ser	Asn	Met	Gly	Phe	Ser	Gly	Asp	Lys
	610					615					620				
Asn	Glu	Leu	Ile	Ile	Gly	Ala	Glu	Ser	Phe	Val	Ser	Asn	Glu	Lys	Ile
625					630					635					640
Tyr	Ile	Asp	Lys	Ile	Glu	Phe	Ile	Pro	Val	Gln	Leu				
				645					650						

<210> 12

<211> 1180

<212> PRT

<213> *Bacillus thuringiensis*

<400> 12

Met	Asn	Pro	Tyr	Gln	Asn	Lys	Asn	Glu	Tyr	Glu	Thr	Leu	Asn	Ala	Ser
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Gln	Lys	Lys	Leu	Asn	Ile	Ser	Asn	Asn	Tyr	Thr	Arg	Tyr	Pro	Ile	Glu
			20					25					30		
Asn	Ser	Pro	Lys	Gln	Leu	Leu	Gln	Ser	Thr	Asn	Tyr	Lys	Asp	Trp	Leu
		35					40					45			
Asn	Met	Cys	Gln	Gln	Asn	Gln	Tyr	Gly	Gly	Asp	Phe	Glu	Thr	Phe	
	50					55				60					
Ile	Asp	Ser	Gly	Glu	Leu	Ser	Ala	Tyr	Thr	Ile	Val	Val	Gly	Thr	Val
65					70					75				80	
Leu	Thr	Gly	Phe	Gly	Phe	Thr	Thr	Pro	Leu	Gly	Leu	Ala	Leu	Ile	Gly
				85					90					95	
Phe	Gly	Thr	Leu	Ile	Pro	Val	Leu	Phe	Pro	Ala	Gln	Asp	Gln	Ser	Asn

His	Thr	Gly	Gly	Asp	Leu	Ile	Asp	Phe	Lys	Asp	His	Phe	Lys	Ile	Thr
				565					570					575	
Cys	Gln	His	Ser	Asn	Phe	Gln	Gln	Ser	Tyr	Phe	Ile	Arg	Ile	Arg	Tyr
			580					585					590		
Ala	Ser	Asn	Gly	Ser	Ala	Asn	Thr	Arg	Ala	Val	Ile	Asn	Leu	Ser	Ile
		595					600					605			
Pro	Gly	Val	Ala	Glu	Leu	Gly	Met	Ala	Leu	Asn	Pro	Thr	Phe	Ser	Gly
	610					615					620				
Thr	Asp	Tyr	Thr	Asn	Leu	Lys	Tyr	Lys	Asp	Phe	Gln	Tyr	Leu	Glu	Phe
625					630					635					640
Ser	Asn	Glu	Val	Lys	Phe	Ala	Pro	Asn	Gln	Asn	Ile	Ser	Leu	Val	Phe
				645					650					655	
Asn	Arg	Ser	Asp	Val	Tyr	Thr	Asn	Thr	Thr	Val	Leu	Ile	Asp	Lys	Ile
			660					665					670		
Glu	Phe	Leu	Pro	Ile	Thr	Arg	Ser	Ile	Arg	Glu	Asp	Arg	Glu	Lys	Gln
		675					680					685			
Lys	Leu	Glu	Thr	Val	Gln	Gln	Ile	Ile	Asn	Thr	Phe	Tyr	Ala	Asn	Pro
	690					695					700				
Ile	Lys	Asn	Thr	Leu	Gln	Ser	Glu	Leu	Thr	Asp	Tyr	Asp	Ile	Asp	Gln
705					710					715					720
Ala	Ala	Asn	Leu	Val	Glu	Cys	Ile	Ser	Glu	Glu	Leu	Tyr	Pro	Lys	Glu
				725					730					735	
Lys	Met	Leu	Leu	Leu	Asp	Glu	Val	Lys	Asn	Ala	Lys	Gln	Leu	Ser	Gln
		740						745				750			
Ser	Arg	Asn	Val	Leu	Gln	Asn	Gly	Asp	Phe	Glu	Ser	Ala	Thr	Leu	Gly
		755					760					765			
Trp	Thr	Thr	Ser	Asp	Asn	Ile	Thr	Ile	Gln	Glu	Asp	Asp	Pro	Ile	Phe
	770					775						780			
Lys	Gly	His	Tyr	Leu	His	Met	Ser	Gly	Ala	Arg	Asp	Ile	Asp	Gly	Thr
785					790					795					800
Ile	Phe	Pro	Thr	Tyr	Ile	Phe	Gln	Lys	Ile	Asp	Glu	Ser	Lys	Leu	Lys
				805					810					815	
Pro	Tyr	Thr	Arg	Tyr	Leu	Val	Arg	Gly	Phe	Val	Gly	Ser	Ser	Lys	Asp
			820					825					830		
Val	Glu	Leu	Val	Val	Ser	Arg	Tyr	Gly	Glu	Glu	Ile	Asp	Ala	Ile	Met
		835					840					845			
Asn	Val	Pro	Ala	Asp	Leu	Asn	Tyr	Leu	Tyr	Pro	Ser	Thr	Phe	Asp	Cys
	850					855					860				
Glu	Gly	Ser	Asn	Arg	Cys	Glu	Thr	Ser	Ala	Val	Pro	Ala	Asn	Ile	Gly
865					870					875					880
Asn	Thr	Ser	Asp	Met	Leu	Tyr	Ser	Cys	Gln	Tyr	Asp	Thr	Gly	Lys	Lys
				885					890					895	
His	Val	Val	Cys	Gln	Asp	Ser	His	Gln	Phe	Ser	Phe	Thr	Ile	Asp	Thr
			900					905					910		
Gly	Ala	Leu	Asp	Thr	Asn	Glu	Asn	Ile	Gly	Val	Trp	Val	Met	Phe	Lys
		915					920					925			
Ile	Ser	Ser	Pro	Asp	Gly	Tyr	Ala	Ser	Leu	Asp	Asn	Leu	Glu	Val	Ile
	930					935					940				
Glu	Glu	Gly	Pro	Ile	Asp	Gly	Glu	Ala	Leu	Ser	Arg	Val	Lys	His	Met
945					950					955					960
Glu	Lys	Lys	Trp	Asn	Asp	Gln	Met	Glu	Ala	Lys	Arg	Ser	Glu	Thr	Gln
				965					970					975	
Gln	Ala	Tyr	Asp	Val	Ala	Lys	Gln	Ala	Ile	Asp	Ala	Leu	Phe	Thr	Asn
			980					985					990		
Val	Gln	Asp	Glu	Ala	Leu	Gln	Phe	Asp	Thr	Thr	Leu	Ala	Gln	Ile	Gln
		995					1000					1005			
Tyr	Ala	Glu	Tyr	Leu	Val	Gln	Ser	Ile	Pro	Tyr	Val	Tyr	Asn	Asp	Trp

1010	1015	1020
Leu Ser Asp Val Pro Gly Met Asn Tyr Asp	Ile Tyr Val Glu Leu Asp	
1025	1030	1035
Ala Arg Val Ala Gln Ala Arg Tyr Leu Tyr Asp Thr Arg Asn Ile Ile		1040
	1045	1050
Lys Asn Gly Asp Phe Thr Gln Gly Val Met Gly Trp His Val Thr Gly		1055
	1060	1065
Asn Ala Asp Val Gln Gln Ile Asp Gly Val Ser Val Leu Val Leu Ser		1070
	1075	1080
Asn Trp Ser Ala Gly Val Ser Gln Asn Val His Leu Gln His Asn His		1085
	1090	1095
Gly Tyr Val Leu Arg Val Ile Ala Lys Lys Glu Gly Pro Gly Asn Gly		1100
1105	1110	1115
Tyr Val Thr Leu Met Asp Cys Glu Glu Asn Gln Glu Lys Leu Thr Phe		1120
	1125	1130
Thr Ser Cys Glu Glu Gly Tyr Ile Thr Lys Thr Val Asp Val Phe Pro		1135
	1140	1145
Asp Thr Asp Arg Val Arg Ile Glu Ile Gly Glu Thr Glu Gly Ser Phe		1150
	1155	1160
Tyr Ile Glu Ser Ile Glu Leu Ile Cys Met Asn Glu		1165
1170	1175	1180

<210> 13

<211> 1136

<212> PRT

<213> *Bacillus thuringiensis*

<400> 13

Met Asn Ser Gly Tyr Pro Leu Ala Asn Asp Leu Gln Gly Ser Met Lys	
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Asn Thr Asn Tyr Lys Asp Trp Leu Ala Met Cys Glu Asn Asn Gln Gln	10
	15
	20
Tyr Gly Val Asn Pro Ala Ala Ile Asn Ser Ser Ser Val Ser Thr Ala	25
	30
	35
Leu Lys Val Ala Gly Ala Ile Leu Lys Phe Val Asn Pro Pro Ala Gly	40
	45
	50
Thr Val Leu Thr Val Leu Ser Ala Val Leu Pro Ile Leu Trp Pro Thr	55
65	60
	70
Asn Thr Pro Thr Pro Glu Arg Val Trp Asn Asp Phe Met Thr Asn Thr	75
	80
	85
Gly Asn Leu Ile Asp Gln Thr Val Thr Ala Tyr Val Arg Thr Asp Ala	90
	95
	100
Asn Ala Lys Met Thr Val Val Lys Asp Tyr Leu Asp Gln Tyr Thr Thr	105
	110
	115
Lys Phe Asn Thr Trp Lys Arg Glu Pro Asn Asn Gln Ser Tyr Arg Thr	120
	125
	130
Ala Val Ile Thr Gln Phe Asn Leu Thr Ser Ala Lys Leu Arg Glu Thr	135
145	140
	150
Ala Val Tyr Phe Ser Asn Leu Val Gly Tyr Glu Leu Leu Leu Leu Pro	155
	160
	165
Ile Tyr Ala Gln Val Ala Asn Phe Asn Leu Leu Leu Ile Arg Asp Gly	170
	175
	180
Leu Ile Asn Ala Gln Glu Trp Ser Leu Ala Arg Ser Ala Gly Asp Gln	185
	190
	195
Leu Tyr Asn Thr Met Val Gln Tyr Thr Lys Glu Tyr Ile Ala His Ser	200
	205
210	215
	220

Ile	Thr	Trp	Tyr	Asn	Lys	Gly	Leu	Asp	Val	Leu	Arg	Asn	Lys	Ser	Asn	225	230	235	240
Gly	Gln	Trp	Ile	Thr	Phe	Asn	Asp	Tyr	Lys	Arg	Glu	Met	Thr	Ile	Gln	245	250	255	
Val	Leu	Asp	Ile	Leu	Ala	Leu	Phe	Ala	Ser	Tyr	Asp	Pro	Arg	Arg	Tyr	260	265	270	
Pro	Ala	Asp	Lys	Ile	Asp	Asn	Thr	Lys	Leu	Ser	Lys	Thr	Glu	Phe	Thr	275	280	285	
Arg	Glu	Ile	Tyr	Thr	Ala	Leu	Val	Glu	Ser	Pro	Ser	Ser	Lys	Ser	Ile	290	295	300	
Ala	Ala	Leu	Glu	Ala	Ala	Leu	Thr	Arg	Asp	Val	His	Leu	Phe	Thr	Trp	305	310	315	320
Leu	Lys	Arg	Val	Asp	Phe	Trp	Thr	Asn	Thr	Ile	Tyr	Gln	Asp	Leu	Arg	325	330	335	
Phe	Leu	Ser	Ala	Asn	Lys	Ile	Gly	Phe	Ser	Tyr	Thr	Asn	Ser	Ser	Ala	340	345	350	
Met	Gln	Glu	Ser	Gly	Ile	Tyr	Gly	Ser	Ser	Gly	Phe	Gly	Ser	Asn	Leu	355	360	365	
Thr	His	Gln	Ile	Gln	Leu	Asn	Ser	Asn	Val	Tyr	Lys	Thr	Ser	Ile	Thr	370	375	380	
Asp	Thr	Ser	Ser	Pro	Ser	Asn	Arg	Val	Thr	Lys	Met	Asp	Phe	Tyr	Lys	385	390	395	400
Ile	Asp	Gly	Thr	Leu	Ala	Ser	Tyr	Asn	Ser	Asn	Ile	Thr	Pro	Thr	Pro	405	410	415	
Glu	Gly	Leu	Arg	Thr	Thr	Phe	Phe	Gly	Phe	Ser	Thr	Asn	Glu	Asn	Thr	420	425	430	
Pro	Asn	Gln	Pro	Thr	Val	Asn	Asp	Tyr	Thr	His	Ile	Leu	Ser	Tyr	Ile	435	440	445	
Lys	Thr	Asp	Val	Ile	Asp	Tyr	Asn	Ser	Asn	Arg	Val	Ser	Phe	Ala	Trp	450	455	460	
Thr	His	Lys	Ile	Val	Asp	Pro	Asn	Asn	Gln	Ile	Tyr	Thr	Asp	Ala	Ile	465	470	475	480
Thr	Gln	Val	Pro	Ala	Val	Lys	Ser	Asn	Phe	Leu	Asn	Ala	Thr	Ala	Lys	485	490	495	
Val	Ile	Lys	Gly	Pro	Gly	His	Thr	Gly	Gly	Asp	Leu	Val	Ala	Leu	Thr	500	505	510	
Ser	Asn	Gly	Thr	Leu	Ser	Gly	Arg	Met	Glu	Ile	Gln	Cys	Lys	Thr	Ser	515	520	525	
Ile	Phe	Asn	Asp	Pro	Thr	Arg	Ser	Tyr	Gly	Leu	Arg	Ile	Arg	Tyr	Ala	530	535	540	
Ala	Asn	Ser	Pro	Ile	Val	Leu	Asn	Val	Ser	Tyr	Val	Leu	Gln	Gly	Val	545	550	555	560
Ser	Arg	Gly	Thr	Thr	Ile	Ser	Thr	Glu	Ser	Thr	Phe	Ser	Arg	Pro	Asn	565	570	575	
Asn	Ile	Ile	Pro	Thr	Asp	Leu	Lys	Tyr	Glu	Glu	Phe	Arg	Tyr	Lys	Asp	580	585	590	
Pro	Phe	Asp	Ala	Ile	Val	Pro	Met	Arg	Leu	Ser	Ser	Asn	Gln	Leu	Ile	595	600	605	
Thr	Ile	Ala	Ile	Gln	Pro	Leu	Asn	Met	Thr	Ser	Asn	Asn	Gln	Val	Ile	610	615	620	
Ile	Asp	Arg	Ile	Glu	Ile	Pro	Ile	Thr	Gln	Ser	Val	Leu	Asp	Glu		625	630	635	640
Thr	Glu	Asn	Gln	Asn	Leu	Glu	Ser	Glu	Arg	Glu	Val	Val	Asn	Ala	Leu	645	650	655	
Phe	Thr	Asn	Asp	Ala	Lys	Asp	Ala	Leu	Asn	Ile	Gly	Thr	Thr	Asp	Tyr	660	665	670	
Asp	Ile	Asp	Gln	Ala	Ala	Asn	Leu	Val	Glu	Cys	Ile	Ser	Glu	Glu	Leu				

		675					680					685				
Tyr	Pro	Lys	Glu	Lys	Met	Leu	Leu	Leu	Asp	Glu	Val	Lys	Asn	Ala	Lys	
	690					695					700					
Gln	Leu	Ser	Gln	Ser	Arg	Asn	Val	Leu	Gln	Asn	Gly	Asp	Phe	Glu	Ser	
705					710					715					720	
Ala	Thr	Leu	Gly	Trp	Thr	Thr	Ser	Asp	Asn	Ile	Thr	Ile	Gln	Glu	Asp	
				725					730					735		
Asp	Pro	Ile	Phe	Lys	Gly	His	Tyr	Leu	His	Met	Ser	Gly	Ala	Arg	Asp	
			740					745					750			
Ile	Asp	Gly	Thr	Ile	Phe	Pro	Thr	Tyr	Ile	Phe	Gln	Lys	Ile	Asp	Glu	
		755					760					765				
Ser	Lys	Leu	Lys	Pro	Tyr	Thr	Arg	Tyr	Leu	Val	Arg	Gly	Phe	Val	Gly	
	770					775					780					
Ser	Ser	Lys	Asp	Val	Glu	Leu	Val	Val	Ser	Arg	Tyr	Gly	Glu	Glu	Ile	
785					790					795					800	
Asp	Ala	Ile	Met	Asn	Val	Pro	Ala	Asp	Leu	Asn	Tyr	Leu	Tyr	Pro	Ser	
				805					810					815		
Thr	Phe	Asp	Cys	Glu	Gly	Ser	Asn	Arg	Cys	Glu	Thr	Ser	Ala	Val	Pro	
			820					825					830			
Ala	Asn	Ile	Gly	Asn	Thr	Ser	Asp	Met	Leu	Tyr	Ser	Cys	Gln	Tyr	Asp	
		835					840					845				
Thr	Gly	Lys	Lys	His	Val	Val	Cys	Gln	Asp	Ser	His	Gln	Phe	Ser	Phe	
	850					855					860					
Thr	Ile	Asp	Thr	Gly	Ala	Leu	Asp	Thr	Asn	Glu	Asn	Ile	Gly	Val	Trp	
865					870					875					880	
Val	Met	Phe	Lys	Ile	Ser	Ser	Pro	Asp	Gly	Tyr	Ala	Ser	Leu	Asp	Asn	
				885					890					895		
Leu	Glu	Val	Ile	Glu	Glu	Gly	Pro	Ile	Asp	Gly	Glu	Ala	Leu	Ser	Arg	
			900					905					910			
Val	Lys	His	Met	Glu	Lys	Lys	Trp	Asn	Asp	Gln	Met	Glu	Ala	Lys	Arg	
		915					920					925				
Ser	Glu	Thr	Gln	Gln	Ala	Tyr	Asp	Val	Ala	Lys	Gln	Ala	Ile	Asp	Ala	
	930					935					940					
Leu	Phe	Thr	Asn	Val	Gln	Asp	Glu	Ala	Leu	Gln	Phe	Asp	Thr	Thr	Leu	
945					950					955					960	
Ala	Gln	Ile	Gln	Tyr	Ala	Glu	Tyr	Leu	Val	Gln	Ser	Ile	Pro	Tyr	Val	
				965					970					975		
Tyr	Asn	Asp	Trp	Leu	Ser	Asp	Val	Pro	Gly	Met	Asn	Tyr	Asp	Ile	Tyr	
			980					985					990			
Val	Glu	Leu	Asp	Ala	Arg	Val	Ala	Gln	Ala	Arg	Tyr	Leu	Tyr	Asp	Thr	
		995					1000					1005				
Arg	Asn	Ile	Ile	Lys	Asn	Gly	Asp	Phe	Thr	Gln	Gly	Val	Met	Gly	Trp	
	1010					1015					1020					
His	Val	Thr	Gly	Asn	Ala	Asp	Val	Gln	Gln	Ile	Asp	Gly	Val	Ser	Val	
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<210> 14
 <211> 475
 <212> PRT
 <213> *Bacillus thuringiensis*

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 Thr Ile Lys Leu Asn Ser Asn Lys Lys Tyr Gly Pro Gly Asp Met Thr
 20 25 30
 Asn Gly Asn Gln Phe Ile Ile Ser Lys Gln Glu Trp Ala Thr Ile Gly
 35 40 45
 Ala Tyr Ile Gln Thr Gly Leu Gly Leu Pro Val Asn Glu Gln Gln Leu
 50 55 60
 Arg Thr His Val Asn Leu Ser Gln Asp Ile Ser Ile Pro Ser Asp Phe
 65 70 75 80
 Ser Gln Leu Tyr Asp Val Tyr Cys Ser Asp Lys Thr Ser Ala Glu Trp
 85 90 95
 Trp Asn Lys Asn Leu Tyr Pro Leu Ile Ile Lys Ser Ala Asn Asp Ile
 100 105 110
 Ala Ser Tyr Gly Phe Lys Val Ala Gly Asp Pro Ser Ile Lys Lys Asp
 115 120 125
 Gly Tyr Phe Lys Lys Leu Gln Asp Glu Leu Asp Asn Ile Val Asp Asn
 130 135 140
 Asn Ser Asp Asp Asp Ala Ile Ala Lys Ala Ile Lys Asp Phe Lys Ala
 145 150 155 160
 Arg Cys Gly Ile Leu Ile Lys Glu Ala Lys Gln Tyr Glu Glu Ala Ala
 165 170 175
 Lys Asn Ile Val Thr Ser Leu Asp Gln Phe Leu His Gly Asp Gln Lys
 180 185 190
 Lys Leu Glu Gly Val Ile Asn Ile Gln Lys Arg Leu Lys Glu Val Gln
 195 200 205
 Thr Ala Leu Asn Gln Ala His Gly Glu Ser Ser Pro Ala His Lys Glu
 210 215 220
 Leu Leu Glu Lys Val Lys Asn Leu Lys Thr Thr Leu Glu Arg Thr Ile
 225 230 235 240
 Lys Ala Glu Gln Asp Leu Glu Lys Lys Val Glu Tyr Ser Phe Leu Leu
 245 250 255
 Gly Pro Leu Leu Gly Phe Val Val Tyr Glu Ile Leu Glu Asn Thr Ala
 260 265 270
 Val Gln His Ile Lys Asn Gln Ile Asp Glu Ile Lys Lys Gln Leu Asp
 275 280 285
 Ser Ala Gln His Asp Leu Asp Arg Asp Val Lys Ile Ile Gly Met Leu
 290 295 300
 Asn Ser Ile Asn Thr Asp Ile Asp Asn Leu Tyr Ser Gln Gly Gln Glu
 305 310 315 320
 Ala Ile Lys Val Phe Gln Lys Leu Gln Gly Ile Trp Ala Thr Ile Gly
 325 330 335
 Ala Gln Ile Glu Asn Leu Arg Thr Thr Ser Leu Gln Glu Val Gln Asp
 340 345 350
 Ser Asp Asp Ala Asp Glu Ile Gln Ile Glu Leu Glu Asp Ala Ser Asp
 355 360 365
 Ala Trp Leu Val Val Ala Gln Glu Ala Arg Asp Phe Thr Leu Asn Ala
 370 375 380
 Tyr Ser Thr Asn Ser Arg Gln Asn Leu Pro Ile Asn Val Ile Ser Asp

385					390					395				400	
Ser	Cys	Asn	Cys	Ser	Thr	Thr	Asn	Met	Thr	Ser	Asn	Gln	Tyr	Ser	Asn
				405					410					415	
Pro	Thr	Thr	Asn	Met	Thr	Ser	Asn	Gln	Tyr	Met	Ile	Ser	His	Glu	Tyr
			420					425					430		
Thr	Ser	Leu	Pro	Asn	Asn	Phe	Met	Leu	Ser	Arg	Asn	Ser	Asn	Leu	Glu
		435					440					445			
Tyr	Lys	Cys	Pro	Glu	Asn	Asn	Phe	Met	Ile	Tyr	Trp	Tyr	Asn	Asn	Ser
	450					455					460				
Asp	Trp	Tyr	Asn	Asn	Ser	Asp	Trp	Tyr	Asn	Asn					
465					470					475					

<210> 15
 <211> 1138
 <212> PRT
 <213> *Bacillus thuringiensis*

<400> 15

Met	Asn	Leu	Asn	Asn	Leu	Asp	Gly	Tyr	Glu	Asp	Ser	Asn	Arg	Thr	Leu
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Asn	Asn	Ser	Leu	Asn	Tyr	Pro	Thr	Gln	Lys	Ala	Leu	Ser	Pro	Ser	Leu
			20					25					30		
Lys	Asn	Met	Asn	Tyr	Gln	Asp	Phe	Leu	Ser	Ile	Thr	Glu	Arg	Glu	Gln
		35					40					45			
Pro	Glu	Ala	Leu	Ala	Ser	Gly	Asn	Thr	Ala	Ile	Asn	Thr	Val	Val	Ser
	50					55					60				
Val	Thr	Gly	Ala	Thr	Leu	Ser	Ala	Leu	Gly	Val	Pro	Gly	Ala	Ser	Phe
65					70					75					80
Ile	Thr	Asn	Phe	Tyr	Leu	Lys	Ile	Ala	Gly	Leu	Leu	Trp	Pro	Glu	Asn
				85					90					95	
Gly	Lys	Ile	Trp	Asp	Glu	Phe	Met	Thr	Glu	Val	Glu	Ala	Leu	Ile	Asp
			100					105					110		
Gln	Lys	Ile	Glu	Glu	Tyr	Val	Arg	Asn	Lys	Ala	Ile	Ala	Glu	Leu	Asp
		115						120				125			
Gly	Leu	Gly	Ser	Ala	Leu	Asp	Lys	Tyr	Gln	Lys	Ala	Leu	Ala	Asp	Trp
	130					135					140				
Leu	Gly	Lys	Gln	Asp	Asp	Pro	Glu	Ala	Ile	Leu	Ser	Val	Ala	Thr	Glu
145					150					155					160
Phe	Arg	Ile	Ile	Asp	Ser	Leu	Phe	Glu	Phe	Ser	Met	Pro	Ser	Phe	Lys
				165					170					175	
Val	Thr	Gly	Tyr	Glu	Ile	Pro	Leu	Leu	Thr	Val	Tyr	Ala	Gln	Ala	Ala
			180						185				190		
Asn	Leu	His	Leu	Ala	Leu	Leu	Arg	Asp	Ser	Thr	Leu	Tyr	Gly	Asp	Lys
		195					200						205		
Trp	Gly	Phe	Thr	Gln	Asn	Asn	Ile	Glu	Glu	Asn	Tyr	Asn	Arg	Gln	Lys
	210					215					220				
Lys	Arg	Ile	Ser	Glu	Tyr	Ser	Asp	His	Cys	Thr	Lys	Trp	Tyr	Asn	Ser
225					230					235					240
Gly	Leu	Ser	Arg	Leu	Asn	Gly	Ser	Thr	Tyr	Glu	Gln	Trp	Ile	Asn	Tyr
				245					250					255	
Asn	Arg	Phe	Arg	Arg	Glu	Met	Ile	Leu	Met	Ala	Leu	Asp	Leu	Val	Ala
			260					265					270		
Val	Phe	Pro	Phe	His	Asp	Pro	Arg	Arg	Tyr	Ser	Met	Glu	Thr	Ser	Thr
		275					280					285			
Gln	Leu	Thr	Arg	Glu	Val	Tyr	Thr	Asp	Pro	Val	Ser	Leu	Ser	Ile	Ser
	290					295					300				

Asn	Pro	Asp	Ile	Gly	Pro	Ser	Phe	Ser	Gln	Met	Glu	Asn	Thr	Ala	Ile	305	310	315	320
Arg	Thr	Pro	His	Leu	Val	Asp	Tyr	Leu	Asp	Glu	Leu	Tyr	Ile	Tyr	Thr	325	330	335	
Ser	Lys	Tyr	Lys	Ala	Phe	Ser	His	Glu	Ile	Gln	Pro	Asp	Leu	Phe	Tyr	340	345	350	
Trp	Ser	Ala	His	Lys	Val	Ser	Phe	Lys	Lys	Ser	Glu	Gln	Ser	Asn	Leu	355	360	365	
Tyr	Thr	Thr	Gly	Ile	Tyr	Gly	Lys	Thr	Ser	Gly	Tyr	Ile	Ser	Ser	Gly	370	375	380	
Ala	Tyr	Ser	Phe	His	Gly	Asn	Asp	Ile	Tyr	Arg	Thr	Leu	Ala	Ala	Pro	385	390	395	400
Ser	Val	Val	Val	Tyr	Pro	Tyr	Thr	Gln	Asn	Tyr	Gly	Val	Glu	Gln	Val	405	410	415	
Glu	Phe	Tyr	Gly	Val	Lys	Gly	His	Val	His	Tyr	Arg	Gly	Asp	Asn	Lys	420	425	430	
Tyr	Asp	Leu	Thr	Tyr	Asp	Ser	Ile	Asp	Gln	Leu	Pro	Pro	Asp	Gly	Glu	435	440	445	
Pro	Ile	His	Glu	Lys	Tyr	Thr	His	Arg	Leu	Cys	His	Ala	Thr	Ala	Ile	450	455	460	
Phe	Lys	Ser	Thr	Pro	Asp	Tyr	Asp	Asn	Ala	Thr	Ile	Pro	Ile	Phe	Ser	465	470	475	480
Trp	Thr	His	Arg	Ser	Ala	Glu	Tyr	Tyr	Asn	Arg	Ile	Tyr	Pro	Asn	Lys	485	490	495	
Ile	Thr	Lys	Ile	Pro	Ala	Val	Lys	Met	Tyr	Lys	Leu	Asp	Asp	Pro	Ser	500	505	510	
Thr	Val	Val	Lys	Gly	Pro	Gly	Phe	Thr	Gly	Gly	Asp	Leu	Val	Lys	Arg	515	520	525	
Gly	Ser	Thr	Gly	Tyr	Ile	Gly	Asp	Ile	Lys	Ala	Thr	Val	Asn	Ser	Pro	530	535	540	
Leu	Ser	Gln	Lys	Tyr	Arg	Val	Arg	Val	Arg	Tyr	Ala	Thr	Asn	Val	Ser	545	550	555	560
Gly	Gln	Phe	Asn	Val	Tyr	Ile	Asn	Asp	Lys	Ile	Thr	Leu	Gln	Thr	Lys	565	570	575	
Phe	Gln	Asn	Thr	Val	Glu	Thr	Ile	Gly	Glu	Gly	Lys	Asp	Leu	Thr	Tyr	580	585	590	
Gly	Ser	Phe	Gly	Tyr	Ile	Glu	Tyr	Ser	Thr	Thr	Ile	Gln	Phe	Pro	Asp	595	600	605	
Glu	His	Pro	Lys	Ile	Thr	Leu	His	Leu	Ser	Asp	Leu	Ser	Asn	Asn	Ser	610	615	620	
Ser	Phe	Tyr	Val	Asp	Ser	Ile	Glu	Phe	Ile	Pro	Val	Asp	Val	Asn	Tyr	625	630	635	640
Ala	Glu	Lys	Glu	Lys	Leu	Glu	Lys	Ala	Gln	Lys	Ala	Val	Asn	Thr	Leu	645	650	655	
Phe	Thr	Glu	Gly	Arg	Asn	Ala	Leu	Gln	Lys	Asp	Val	Thr	Asp	Tyr	Lys	660	665	670	
Val	Asp	Gln	Val	Ser	Ile	Leu	Val	Asp	Cys	Ile	Ser	Gly	Asp	Leu	Tyr	675	680	685	
Pro	Asn	Glu	Lys	Arg	Glu	Leu	Gln	Asn	Leu	Val	Lys	Tyr	Ala	Lys	Arg	690	695	700	
Leu	Ser	Tyr	Ser	Arg	Asn	Leu	Leu	Leu	Asp	Pro	Thr	Phe	Asp	Ser	Ile	705	710	715	720
Asn	Ser	Ser	Glu	Glu	Asn	Gly	Trp	Tyr	Gly	Ser	Asn	Gly	Ile	Val	Ile	725	730	735	
Gly	Asn	Gly	Asp	Phe	Val	Phe	Lys	Gly	Asn	Tyr	Leu	Ile	Phe	Ser	Gly	740	745	750	
Thr	Asn	Asp	Thr	Gln	Tyr	Pro	Thr	Tyr	Leu	Tyr	Gln	Lys	Ile	Asp	Glu				

Met	Ser	Pro	Asn	Asn	Gln	Asn	Glu	Tyr	Glu	Ile	Ile	Asp	Ala	Thr	Pro	
1				5					10					15		
Ser	Thr	Ser	Val	Ser	Ser	Asp	Ser	Asn	Arg	Tyr	Pro	Phe	Ala	Asn	Glu	
			20					25					30			
Pro	Thr	Asp	Ala	Leu	Gln	Asn	Met	Asn	Tyr	Lys	Asp	Tyr	Leu	Lys	Met	
		35					40					45				
Ser	Gly	Gly	Glu	Asn	Pro	Glu	Leu	Phe	Gly	Asn	Pro	Glu	Thr	Phe	Ile	
50						55					60					
Ser	Ser	Ser	Thr	Ile	Gln	Thr	Gly	Ile	Gly	Ile	Val	Gly	Arg	Ile	Leu	
65					70					75					80	
Gly	Ala	Leu	Gly	Val	Pro	Phe	Ala	Ser	Gln	Ile	Ala	Ser	Phe	Tyr	Ser	
				85					90					95		
Phe	Ile	Val	Gly	Gln	Leu	Trp	Pro	Ser	Lys	Ser	Val	Asp	Ile	Trp	Gly	
			100					105					110			
Glu	Ile	Met	Glu	Arg	Val	Glu	Glu	Leu	Val	Asp	Gln	Lys	Ile	Glu	Lys	
		115					120					125				
Tyr	Val	Lys	Asp	Lys	Ala	Leu	Ala	Glu	Leu	Lys	Gly	Leu	Gly	Asn	Ala	
130						135						140				
Leu	Asp	Val	Tyr	Gln	Gln	Ser	Leu	Glu	Asp	Trp	Leu	Glu	Asn	Arg	Asn	
145					150					155					160	
Asp	Ala	Arg	Thr	Arg	Ser	Val	Val	Ser	Asn	Gln	Phe	Ile	Ala	Leu	Asp	
				165					170					175		
Leu	Asn	Phe	Val	Ser	Ser	Ile	Pro	Ser	Phe	Ala	Val	Ser	Gly	His	Glu	
			180					185					190			
Val	Leu	Leu	Leu	Ala	Val	Tyr	Ala	Gln	Ala	Val	Asn	Leu	His	Leu	Leu	
		195					200					205				
Leu	Leu	Arg	Asp	Ala	Ser	Ile	Phe	Gly	Glu	Glu	Trp	Gly	Phe	Thr	Pro	
210						215						220				
Gly	Glu	Ile	Ser	Arg	Phe	Tyr	Asn	Arg	Gln	Val	Gln	Leu	Thr	Ala	Glu	
225					230					235					240	
Tyr	Ser	Asp	Tyr	Cys	Val	Lys	Trp	Tyr	Lys	Ile	Gly	Leu	Asp	Lys	Leu	
				245					250					255		
Lys	Gly	Thr	Thr	Ser	Lys	Ser	Trp	Leu	Asn	Tyr	His	Gln	Phe	Arg	Arg	
			260					265						270		
Glu	Met	Thr	Leu	Leu	Val	Leu	Asp	Leu	Val	Ala	Leu	Phe	Pro	Asn	Tyr	
		275					280					285				
Asp	Thr	His	Met	Tyr	Pro	Ile	Glu	Thr	Thr	Ala	Gln	Leu	Thr	Arg	Asp	
290						295					300					
Val	Tyr	Thr	Asp	Pro	Ile	Ala	Phe	Asn	Ile	Val	Thr	Ser	Thr	Gly	Phe	
305					310					315					320	
Cys	Asn	Pro	Trp	Ser	Thr	His	Ser	Gly	Ile	Leu	Phe	Tyr	Glu	Val	Glu	
				325					330					335		
Asn	Asn	Val	Ile	Arg	Pro	Pro	His	Leu	Phe	Asp	Ile	Leu	Ser	Ser	Val	
			340					345					350			
Glu	Ile	Asn	Thr	Ser	Arg	Gly	Gly	Ile	Thr	Leu	Asn	Asn	Asp	Ala	Tyr	
		355					360					365				
Ile	Asn	Tyr	Trp	Ser	Gly	His	Thr	Leu	Lys	Tyr	Arg	Arg	Thr	Ala	Asp	
370						375						380				
Ser	Thr	Val	Thr	Tyr	Thr	Ala	Asn	Tyr	Gly	Arg	Ile	Thr	Ser	Glu	Lys	
385					390					395					400	
Asn	Ser	Phe	Ala	Leu	Glu	Asp	Arg	Asp	Ile	Phe	Glu	Ile	Asn	Ser	Thr	
				405					410					415		
Val	Ala	Asn	Leu	Ala	Asn	Tyr	Tyr	Gln	Lys	Ala	Tyr	Gly	Val	Pro	Gly	
			420					425					430			
Ser	Trp	Phe	His	Met	Val	Lys	Arg	Gly	Thr	Ser	Ser	Thr	Thr	Ala	Tyr	
		435					440					445				
Leu	Tyr	Ser	Lys	Thr	His	Thr	Ala	Leu	Gln	Gly	Cys	Thr	Gln	Val	Tyr	

450		455		460
Glu Ser Ser Asp Glu Ile Pro Leu Asp Arg Thr Val Pro Val Ala Glu				
465		470		475
Ser Tyr Ser His Arg Leu Ser His Ile Thr Ser His Ser Phe Ser Lys				
	485		490	495
Asn Gly Ser Ala Tyr Tyr Gly Ser Phe Pro Val Phe Val Trp Thr His				
	500		505	510
Thr Ser Ala Asp Leu Asn Asn Thr Ile Tyr Ser Asp Lys Ile Thr Gln				
	515		520	525
Ile Pro Ala Val Lys Gly Asp Met Leu Tyr Leu Gly Gly Ser Val Val				
	530		535	540
Gln Gly Pro Gly Phe Thr Gly Gly Asp Ile Leu Lys Arg Thr Asn Pro				
545		550		555
Ser Ile Leu Gly Thr Phe Ala Val Thr Val Asn Gly Ser Leu Ser Gln				
	565		570	575
Arg Tyr Arg Val Arg Ile Arg Tyr Ala Ser Thr Thr Asp Phe Glu Phe				
	580		585	590
Thr Leu Tyr Leu Gly Asp Thr Ile Glu Lys Asn Arg Phe Asn Lys Thr				
	595		600	605
Met Asp Asn Gly Ala Ser Leu Thr Tyr Glu Thr Phe Lys Phe Ala Ser				
	610		615	620
Phe Ile Thr Asp Phe Gln Phe Arg Glu Thr Gln Asp Lys Ile Leu Leu				
625		630		635
Ser Met Gly Asp Phe Ser Ser Gly Gln Glu Val Tyr Ile Asp Arg Ile				
	645		650	655
Glu Phe Ile Pro Val Asp Glu Thr Tyr Glu Ala Glu Gln Asp Leu Glu				
	660		665	670
Ala Ala Lys Lys Ala Val Asn Ala Leu Phe Thr Asn Thr Lys Asp Gly				
	675		680	685
Leu Arg Pro Gly Val Thr Asp Tyr Glu Val Asn Gln Ala Ala Asn Leu				
	690		695	700
Val Glu Cys Leu Ser Asp Leu Tyr Pro Asn Glu Lys Arg Leu Leu				
705		710		715
Phe Asp Ala Val Arg Glu Ala Lys Arg Leu Ser Gly Ala Arg Asn Leu				
	725		730	735
Leu Gln Asp Pro Asp Phe Gln Glu Ile Asn Gly Glu Asn Gly Trp Ala				
	740		745	750
Ala Ser Thr Gly Ile Glu Ile Val Glu Gly Asp Ala Val Phe Lys Gly				
	755		760	765
Arg Tyr Leu Arg Leu Pro Gly Ala Arg Glu Ile Asp Thr Glu Thr Tyr				
	770		775	780
Pro Thr Tyr Leu Tyr Gln Lys Val Glu Glu Gly Val Leu Lys Pro Tyr				
785		790		795
Thr Arg Tyr Arg Leu Arg Gly Phe Val Gly Ser Ser Gln Gly Leu Glu				
	805		810	815
Ile Tyr Thr Ile Arg His Gln Thr Asn Arg Ile Val Lys Asn Val Pro				
	820		825	830
Asp Asp Leu Leu Pro Asp Val Ser Pro Val Asn Ser Asp Gly Ser Ile				
	835		840	845
Asn Arg Cys Ser Glu Gln Lys Tyr Val Asn Ser Arg Leu Glu Gly Glu				
	850		855	860
Asn Arg Ser Gly Asp Ala His Glu Phe Ser Leu Pro Ile Asp Ile Gly				
865		870		875
Glu Leu Asp Tyr Asn Glu Asn Ala Gly Ile Trp Val Gly Phe Lys Ile				
	885		890	895
Thr Asp Pro Glu Gly Tyr Ala Thr Leu Gly Asn Leu Glu Leu Val Glu				
	900		905	910

Glu Gly Pro Leu Ser Gly Asp Ala Leu Glu Arg Leu Gln Arg Glu Glu
 915 920 925
 Gln Gln Trp Lys Ile Gln Met Thr Arg Arg Arg Glu Glu Thr Asp Arg
 930 935 940
 Arg Tyr Met Ala Ser Lys Gln Ala Val Asp Arg Leu Tyr Ala Asp Tyr
 945 950 955 960
 Gln Asp Gln Gln Leu Asn Pro Asp Val Glu Ile Thr Asp Leu Thr Ala
 965 970 975
 Ala Gln Asp Leu Ile Gln Ser Ile Pro Tyr Val Tyr Asn Glu Met Phe
 980 985 990
 Pro Glu Ile Pro Gly Met Asn Tyr Thr Lys Phe Thr Glu Leu Thr Asp
 995 1000 1005
 Arg Leu Gln Gln Ala Trp Asn Leu Tyr Asp Gln Arg Asn Ala Ile Pro
 1010 1015 1020
 Asn Gly Asp Phe Arg Asn Gly Leu Ser Asn Trp Asn Ala Thr Pro Gly
 1025 1030 1035 1040
 Val Glu Val Gln Gln Ile Asn His Thr Ser Val Leu Val Ile Pro Asn
 1045 1050 1055
 Trp Asp Glu Gln Val Ser Gln Gln Phe Thr Val Gln Pro Asn Gln Arg
 1060 1065 1070
 Tyr Val Leu Arg Val Thr Ala Arg Lys Glu Gly Val Gly Asn Gly Tyr
 1075 1080 1085
 Val Ser Ile Arg Asp Gly Gly Asn Gln Ser Glu Thr Leu Thr Phe Ser
 1090 1095 1100
 Ala Ser Asp Tyr Asp Thr Asn Gly Val Tyr Asn Asp Gln Thr Gly Tyr
 1105 1110 1115 1120
 Ile Thr Lys Thr Val Thr Phe Ile Pro Tyr Thr Asp Gln Met Trp Ile
 1125 1130 1135
 Glu Ile Ser Glu Thr Glu Gly Thr Phe Tyr Ile Glu Ser Val Glu Leu
 1140 1145 1150
 Ile Val Asp Val Glu
 1155

<210> 17
 <211> 675
 <212> PRT
 <213> *Bacillus thuringiensis*

<400> 17
 Met Asn Pro Tyr Gln Asn Lys Asn Glu Tyr Glu Ile Phe Asn Ala Pro
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 Ser Asn Gly Phe Ser Lys Ser Asn Asn Tyr Ser Arg Tyr Pro Leu Ala
 20 25 30
 Asn Lys Pro Asn Gln Pro Leu Lys Asn Thr Asn Tyr Lys Asp Trp Leu
 35 40 45
 Asn Val Cys Gln Asp Asn Gln Gln Tyr Gly Asn Asn Ala Gly Asn Phe
 50 55 60
 Ala Ser Ser Glu Thr Ile Val Gly Val Ser Ala Gly Ile Ile Val Val
 65 70 75 80
 Gly Thr Met Leu Gly Ala Phe Ala Ala Pro Val Leu Ala Ala Gly Ile
 85 90 95
 Ile Ser Phe Gly Thr Leu Leu Pro Ile Phe Trp Gln Gly Ser Asp Pro
 100 105 110
 Ala Asn Val Trp Gln Asp Leu Leu Asn Ile Gly Gly Arg Pro Ile Gln
 115 120 125
 Glu Ile Asp Lys Asn Ile Ile Asn Val Leu Thr Ser Ile Val Thr Pro

130		135		140
Ile Lys Asn Gln Leu Asp Lys Tyr Gln Glu Phe Phe Asp Lys Trp Glu				
145		150		160
Pro Ala Arg Thr His Ala Asn Ala Lys Ala Val His Asp Leu Phe Thr				
	165		170	175
Thr Leu Glu Pro Ile Ile Asp Lys Asp Leu Asp Met Leu Lys Asn Asn				
	180		185	190
Ala Ser Tyr Arg Ile Pro Thr Leu Pro Ala Tyr Ala Gln Ile Ala Thr				
	195		200	205
Trp His Leu Asn Leu Leu Lys His Ala Ala Thr Tyr Tyr Asn Ile Trp				
	210		215	220
Leu Gln Asn Gln Gly Ile Asn Pro Ser Thr Phe Asn Ser Ser Asn Tyr				
225		230		240
Tyr Gln Gly Tyr Leu Lys Arg Lys Ile Gln Glu Tyr Thr Asp Tyr Cys				
	245		250	255
Ile Gln Thr Tyr Asn Ala Gly Leu Thr Met Ile Arg Thr Asn Thr Asn				
	260		265	270
Ala Thr Trp Asn Met Tyr Asn Thr Tyr Arg Leu Glu Met Thr Leu Thr				
	275		280	285
Val Leu Asp Leu Ile Ala Ile Phe Pro Asn Tyr Asp Pro Glu Lys Tyr				
	290		295	300
Pro Ile Gly Val Lys Ser Glu Leu Ile Arg Glu Val Tyr Thr Asn Val				
305		310		320
Asn Ser Asp Thr Phe Arg Thr Ile Thr Glu Leu Glu Asn Gly Leu Thr				
	325		330	335
Arg Asn Pro Thr Leu Phe Thr Trp Ile Asn Gln Gly Arg Phe Tyr Thr				
	340		345	350
Arg Asn Ser Arg Asp Ile Leu Asp Pro Tyr Asp Ile Phe Ser Phe Thr				
	355		360	365
Gly Asn Gln Met Ala Phe Thr His Thr Asn Asp Asp Arg Asn Ile Ile				
	370		375	380
Trp Gly Ala Val His Gly Asn Ile Ile Ser Gln Asp Thr Ser Lys Val				
385		390		400
Phe Pro Phe Tyr Arg Asn Lys Pro Ile Asp Lys Val Glu Ile Val Arg				
	405		410	415
His Arg Glu Tyr Ser Asp Ile Ile Tyr Glu Met Ile Phe Phe Ser Asn				
	420		425	430
Ser Ser Glu Val Phe Arg Tyr Ser Ser Asn Ser Thr Ile Glu Asn Asn				
	435		440	445
Tyr Lys Arg Thr Asp Ser Tyr Met Ile Pro Lys Gln Thr Trp Lys Asn				
	450		455	460
Glu Glu Tyr Gly His Thr Leu Ser Tyr Ile Lys Thr Asp Asn Tyr Ile				
465		470		480
Phe Ser Val Val Arg Glu Arg Arg Arg Val Ala Phe Ser Trp Thr His				
	485		490	495
Thr Ser Val Asp Phe Gln Asn Thr Ile Asp Leu Asp Asn Ile Thr Gln				
	500		505	510
Ile His Ala Leu Lys Ala Leu Lys Val Ser Ser Asp Ser Lys Ile Val				
	515		520	525
Lys Gly Pro Gly His Thr Gly Gly Asp Leu Val Ile Leu Lys Asp Ser				
	530		535	540
Met Asp Phe Arg Val Arg Phe Leu Lys Asn Val Ser Arg Gln Tyr Gln				
545		550		560
Val Arg Ile Arg Tyr Ala Thr Asn Ala Pro Lys Thr Thr Val Phe Leu				
	565		570	575
Thr Gly Ile Asp Thr Ile Ser Val Glu Leu Pro Ser Thr Thr Ser Arg				
	580		585	590

Gln	Asn	Pro	Asn	Ala	Thr	Asp	Leu	Thr	Tyr	Ala	Asp	Phe	Gly	Tyr	Val
		595					600					605			
Thr	Phe	Pro	Arg	Thr	Val	Pro	Asn	Lys	Thr	Phe	Glu	Gly	Glu	Asp	Thr
	610					615				620					
Leu	Leu	Met	Thr	Leu	Tyr	Gly	Thr	Pro	Asn	His	Ser	Tyr	Asn	Ile	Tyr
625					630					635					640
Ile	Asp	Lys	Ile	Glu	Phe	Ile	Pro	Ile	Thr	Gln	Ser	Val	Leu	Asp	Tyr
				645					650					655	
Thr	Glu	Lys	Gln	Asn	Ile	Glu	Lys	Thr	Gln	Lys	Ile	Val	Asn	Asp	Leu
			660					665					670		
Phe	Val	Asn													
		675													

<210> 18

<211> 648

<212> PRT

<213> Bacillus thuringiensis

<400> 18

Met	His	Tyr	Tyr	Gly	Asn	Arg	Asn	Glu	Tyr	Asp	Ile	Leu	Asn	Ala	Ser
1				5				10					15		
Ser	Asn	Asp	Ser	Asn	Met	Ser	Asn	Thr	Tyr	Pro	Arg	Tyr	Pro	Leu	Ala
			20				25					30			
Asn	Pro	Gln	Gln	Asp	Leu	Met	Gln	Asn	Thr	Asn	Tyr	Lys	Asp	Trp	Leu
		35				40					45				
Asn	Val	Cys	Glu	Gly	Tyr	His	Ile	Glu	Asn	Pro	Arg	Glu	Ala	Ser	Val
	50					55				60					
Arg	Ala	Gly	Leu	Gly	Lys	Gly	Leu	Gly	Ile	Val	Ser	Thr	Ile	Val	Gly
65					70					75					80
Phe	Phe	Gly	Gly	Ser	Ile	Ile	Leu	Asp	Thr	Ile	Gly	Leu	Phe	Tyr	Gln
				85				90					95		
Ile	Ser	Glu	Leu	Leu	Trp	Pro	Glu	Asp	Asp	Thr	Gln	Gln	Tyr	Thr	Trp
			100					105					110		
Gln	Asp	Ile	Met	Asn	His	Val	Glu	Asp	Leu	Ile	Asp	Lys	Arg	Ile	Thr
		115					120					125			
Glu	Val	Ile	Arg	Gly	Asn	Ala	Ile	Arg	Thr	Leu	Ala	Asp	Leu	Gln	Gly
	130					135					140				
Lys	Val	Asp	Asp	Tyr	Asn	Asn	Trp	Leu	Lys	Lys	Trp	Lys	Asp	Asp	Pro
145					150					155					160
Lys	Ser	Thr	Gly	Asn	Leu	Ser	Thr	Leu	Val	Thr	Lys	Phe	Thr	Ala	Leu
				165				170						175	
Asp	Ser	Asp	Phe	Asn	Gly	Ala	Ile	Arg	Thr	Val	Asn	Asn	Gln	Gly	Ser
			180					185					190		
Pro	Gly	Tyr	Glu	Leu	Leu	Leu	Leu	Pro	Val	Tyr	Ala	Gln	Ile	Ala	Asn
		195					200					205			
Leu	His	Leu	Leu	Leu	Leu	Arg	Asp	Ala	Gln	Ile	Tyr	Gly	Asp	Lys	Trp
	210					215					220				
Trp	Ser	Ala	Arg	Ala	Asn	Ala	Arg	Asp	Asn	Tyr	Tyr	Gln	Ile	Gln	Leu
225					230					235					240
Glu	Lys	Thr	Lys	Glu	Tyr	Thr	Glu	Tyr	Cys	Ile	Asn	Trp	Tyr	Asn	Lys
				245					250					255	
Gly	Leu	Asn	Asp	Phe	Arg	Thr	Ala	Gly	Gln	Trp	Val	Asn	Phe	Asn	Arg
			260					265					270		
Tyr	Arg	Arg	Glu	Met	Thr	Leu	Thr	Val	Leu	Asp	Ile	Ile	Ser	Met	Phe
		275					280					285			
Pro	Ile	Tyr	Asp	Ala	Arg	Leu	Tyr	Pro	Thr	Glu	Val	Lys	Thr	Glu	Leu

290		295		300
Thr Arg Glu Ile Tyr Ser Asp Val Ile Asn Gly Glu Ile Tyr Gly Leu				
305		310		315
Met Thr Pro Tyr Phe Ser Phe Glu Lys Ala Glu Ser Leu Tyr Thr Arg				
	325		330	
Ala Pro His Leu Phe Thr Trp Leu Lys Gly Phe Arg Phe Val Thr Asn				
	340		345	
Ser Ile Ser Tyr Trp Thr Phe Leu Ser Gly Gly Gln Asn Lys Tyr Ser				
	355		360	
Tyr Thr Asn Asn Ser Ser Ile Asn Glu Gly Ser Phe Arg Gly Gln Asp				
	370		375	
Thr Asp Tyr Gly Gly Thr Ser Ser Thr Ile Asn Ile Pro Ser Asn Ser				
385		390		395
Tyr Val Tyr Asn Leu Trp Thr Glu Asn Tyr Glu Tyr Ile Tyr Pro Trp				
	405		410	
Gly Asp Pro Val Asn Ile Thr Lys Met Asn Phe Ser Val Thr Asp Asn				
	420		425	
Asn Ser Ser Lys Glu Leu Ile Tyr Gly Ala His Arg Thr Asn Lys Pro				
	435		440	
Val Val Arg Thr Asp Phe Asp Phe Leu Thr Asn Lys Glu Gly Thr Glu				
	450		455	
Leu Ala Lys Tyr Asn Asp Tyr Asn His Ile Leu Ser Tyr Met Leu Ile				
465		470		475
Asn Gly Glu Thr Phe Gly Gln Lys Arg His Gly Tyr Ser Phe Ala Phe				
	485		490	
Thr His Ser Ser Val Asp Pro Asn Asn Thr Ile Ala Ala Asn Lys Ile				
	500		505	
Thr Gln Ile Pro Val Val Lys Ala Ser Ser Ile Asn Gly Ser Ile Ser				
	515		520	
Ile Glu Lys Gly Pro Gly Phe Thr Gly Gly Asp Leu Val Lys Met Arg				
	530		535	
Ala Asp Ser Gly Leu Thr Met Arg Phe Lys Ala Glu Leu Leu Asp Lys				
545		550		555
Lys Tyr Arg Val Arg Ile Arg Tyr Lys Cys Asn Tyr Ser Ser Lys Leu				
	565		570	
Ile Leu Arg Lys Trp Lys Gly Glu Gly Tyr Ile Gln Gln Gln Ile His				
	580		585	
Asn Ile Ser Pro Thr Tyr Gly Ala Phe Ser Tyr Leu Glu Ser Phe Thr				
	595		600	
Ile Thr Thr Thr Glu Asn Ile Phe Asp Leu Thr Met Glu Val Thr Tyr				
	610		615	
Pro Tyr Gly Arg Gln Phe Val Glu Asp Ile Pro Ser Leu Ile Leu Asp				
625		630		635
Lys Ile Glu Phe Leu Pro Thr Asn				
	645			

<210> 19

<211> 722

<212> PRT

<213> Bacillus thuringiensis

<400> 19

Met Lys Glu Gln Asn Leu Asn Lys Tyr Asp Glu Ile Thr Val Gln Ala
1 5 10 15
Ala Ser Asp Tyr Ile Asp Ile Arg Pro Ile Phe Gln Thr Asn Gly Ser
20 25 30

Ala	Thr	Phe	Asn	Ser	Asn	Thr	Asn	Ile	Thr	Thr	Leu	Thr	Gln	Ala	Ile	
		35					40					45				
Asn	Ser	Gln	Ala	Gly	Ala	Ile	Ala	Gly	Lys	Thr	Ala	Leu	Asp	Met	Arg	
		50				55					60					
His	Asp	Phe	Thr	Phe	Arg	Ala	Asp	Ile	Phe	Leu	Gly	Thr	Lys	Ser	Asn	
65					70					75					80	
Gly	Ala	Asp	Gly	Ile	Ala	Ile	Ala	Phe	His	Arg	Gly	Ser	Ile	Gly	Phe	
				85					90					95		
Val	Gly	Thr	Lys	Gly	Gly	Gly	Leu	Gly	Ile	Leu	Gly	Ala	Pro	Lys	Gly	
			100					105					110			
Ile	Gly	Phe	Glu	Leu	Asp	Thr	Tyr	Ala	Asn	Ala	Pro	Glu	Asp	Glu	Val	
		115					120					125				
Gly	Asp	Ser	Phe	Gly	His	Gly	Ala	Met	Lys	Gly	Ser	Phe	Pro	Ser	Phe	
		130				135					140					
Pro	Asn	Gly	Tyr	Pro	His	Ala	Gly	Phe	Val	Ser	Thr	Asp	Lys	Asn	Ser	
145					150					155					160	
Arg	Trp	Leu	Ser	Ala	Leu	Ala	Gln	Met	Gln	Arg	Ile	Ala	Ala	Pro	Asn	
				165					170						175	
Gly	Arg	Trp	Arg	Arg	Leu	Glu	Ile	Arg	Trp	Asp	Ala	Arg	Asn	Lys	Glu	
			180					185					190			
Leu	Thr	Ala	Asn	Leu	Gln	Asp	Leu	Thr	Phe	Asn	Asp	Ile	Thr	Val	Gly	
		195					200					205				
Glu	Lys	Pro	Arg	Thr	Pro	Arg	Thr	Ala	Thr	Trp	Arg	Leu	Val	Asn	Pro	
		210				215					220					
Ala	Phe	Glu	Leu	Asp	Gln	Lys	Tyr	Thr	Phe	Val	Ile	Gly	Ser	Ala	Thr	
225					230					235					240	
Gly	Ala	Ser	Asn	Asn	Leu	His	Gln	Ile	Gly	Ile	Ile	Glu	Phe	Asp	Ala	
				245					250					255		
Tyr	Phe	Thr	Lys	Pro	Thr	Ile	Glu	Ala	Asn	Asn	Val	Asn	Val	Pro	Val	
			260					265					270			
Gly	Ala	Thr	Phe	Asn	Pro	Lys	Thr	Tyr	Pro	Gly	Ile	Asn	Leu	Arg	Ala	
		275					280					285				
Thr	Asp	Glu	Ile	Asp	Gly	Asp	Leu	Thr	Ser	Lys	Ile	Ile	Val	Lys	Ala	
		290				295					300					
Asn	Asn	Val	Asn	Thr	Ser	Lys	Thr	Gly	Val	Tyr	Tyr	Val	Thr	Tyr	Tyr	
305					310					315					320	
Val	Glu	Asn	Ser	Tyr	Gly	Glu	Ser	Asp	Glu	Lys	Thr	Ile	Glu	Val	Thr	
				325					330					335		
Val	Phe	Ser	Asn	Pro	Thr	Ile	Ile	Ala	Ser	Asp	Val	Glu	Ile	Glu	Lys	
			340					345					350			
Gly	Glu	Ser	Phe	Asn	Pro	Leu	Thr	Asp	Ser	Arg	Val	Gly	Leu	Ser	Ala	
		355					360					365				
Gln	Asp	Ser	Leu	Gly	Asn	Asp	Ile	Thr	Gln	Asn	Val	Lys	Val	Lys	Ser	
		370				375					380					
Ser	Asn	Val	Asp	Thr	Ser	Lys	Pro	Gly	Glu	Tyr	Glu	Val	Val	Phe	Glu	
385					390					395					400	
Val	Thr	Asp	Ser	Phe	Gly	Gly	Lys	Ala	Glu	Lys	Asp	Phe	Lys	Val	Thr	
				405					410					415		
Val	Leu	Gly	Gln	Pro	Ser	Ile	Glu	Ala	Asn	Asn	Val	Glu	Leu	Glu	Ile	
			420					425					430			
Asp	Asp	Ser	Leu	Asp	Pro	Leu	Thr	Asp	Ala	Lys	Val	Gly	Leu	Arg	Ala	
		435					440					445				
Lys	Asp	Ser	Leu	Gly	Asn	Asp	Ile	Thr	Lys	Asp	Ile	Lys	Val	Lys	Phe	
		450				455					460					
Asn	Asn	Val	Asp	Thr	Ser	Asn	Ser	Gly	Lys	Tyr	Glu	Val	Ile	Phe	Glu	
465					470					475					480	
Val	Thr	Asp	Arg	Phe	Gly	Lys	Lys	Ala	Glu	Lys	Ser	Ile	Glu	Val	Leu	

Ala	Trp	Asn	Asn	Asn	Lys	Ser	Asn	Ile	Asn	Tyr	Gln	Thr	Asn	Val	Ala
145					150					155					160
Glu	Ala	Phe	Lys	Thr	Val	Glu	Arg	Glu	Phe	Phe	Thr	Lys	Leu	Lys	Gly
				165						170					175
Ile	Tyr	Arg	Thr	Ser	Ser	Ser	Gln	Ile	Thr	Leu	Leu	Pro	Thr	Phe	Thr
			180						185				190		
Gln	Ala	Ala	Asn	Leu	His	Leu	Ser	Met	Leu	Arg	Asp	Ala	Val	Met	Tyr
		195						200				205			
Gln	Glu	Gly	Trp	Asn	Leu	Gln	Ser	His	Ile	Asn	Tyr	Ser	Lys	Glu	Leu
	210					215					220				
Asp	Asp	Ala	Leu	Glu	Asp	Tyr	Thr	Asn	Tyr	Cys	Val	Glu	Val	Tyr	Thr
225					230					235					240
Lys	Gly	Leu	Asn	Ala	Leu	Arg	Gly	Ser	Thr	Ala	Ile	Asp	Trp	Leu	Glu
			245						250					255	
Phe	Asn	Ser	Phe	Arg	Arg	Asp	Met	Thr	Leu	Met	Val	Leu	Asp	Leu	Val
			260					265					270		
Ala	Ile	Phe	Pro	Asn	Tyr	Asn	Pro	Val	Arg	Tyr	Pro	Leu	Ser	Thr	Lys
		275					280					285			
Ile	Ser	Leu	Ser	Arg	Lys	Ile	Tyr	Thr	Asp	Pro	Val	Gly	Arg	Thr	Asp
	290					295					300				
Ser	Pro	Ser	Phe	Gly	Asp	Trp	Thr	Asn	Thr	Gly	Arg	Thr	Leu	Ala	Asn
305					310					315					320
Phe	Asn	Asp	Leu	Glu	Arg	Glu	Val	Thr	Asp	Ser	Pro	Ser	Leu	Val	Lys
			325						330					335	
Trp	Leu	Gly	Asp	Met	Thr	Ile	Tyr	Thr	Gly	Ala	Ile	Asp	Ser	Tyr	Arg
			340					345					350		
Pro	Thr	Ser	Pro	Gly	Asp	Arg	Ile	Gly	Val	Trp	Tyr	Gly	Asn	Ile	Asn
		355					360					365			
Ala	Phe	Tyr	His	Thr	Gly	Arg	Thr	Asp	Val	Val	Met	Phe	Arg	Gln	Thr
	370					375					380				
Gly	Asp	Thr	Ala	Tyr	Glu	Asp	Pro	Ser	Thr	Phe	Ile	Ser	Asn	Ile	Leu
385					390					395					400
Tyr	Asp	Asp	Ile	Tyr	Lys	Leu	Asp	Leu	Arg	Ala	Ala	Ala	Val	Ser	Thr
			405						410					415	
Ile	Gln	Gly	Ala	Met	Asp	Thr	Thr	Phe	Gly	Val	Ser	Ser	Ser	Arg	Phe
			420					425					430		
Phe	Asp	Ile	Arg	Gly	Arg	Asn	Gln	Leu	Tyr	Gln	Ser	Asn	Lys	Pro	Tyr
		435					440					445			
Pro	Ser	Leu	Pro	Ile	Thr	Ile	Thr	Phe	Pro	Gly	Glu	Glu	Ser	Ser	Glu
	450					455					460				
Gly	Asn	Ala	Asn	Asp	Tyr	Ser	His	Leu	Leu	Cys	Asp	Val	Lys	Ile	Leu
465					470					475					480
Gln	Glu	Asp	Ser	Ser	Asn	Ile	Cys	Glu	Gly	Arg	Ser	Ser	Leu	Leu	Ser
			485						490					495	
His	Ala	Trp	Thr	His	Ala	Ser	Leu	Asp	Arg	Asn	Asn	Thr	Ile	Leu	Pro
			500					505					510		
Asp	Glu	Ile	Thr	Gln	Ile	Pro	Ala	Val	Thr	Ala	Tyr	Glu	Leu	Arg	Gly
		515					520					525			
Asn	Ser	Ser	Val	Val	Ala	Gly	Pro	Gly	Ser	Thr	Gly	Gly	Asp	Leu	Val
	530					535					540				
Lys	Met	Ser	Tyr	His	Ser	Val	Trp	Ser	Phe	Lys	Val	Tyr	Cys	Ser	Glu
545					550					555					560
Leu	Lys	Asn	Tyr	Arg	Val	Arg	Ile	Arg	Tyr	Ala	Ser	His	Gly	Asn	Cys
			565						570					575	
Gln	Phe	Leu	Met	Lys	Arg	Trp	Pro	Ser	Thr	Gly	Val	Ala	Pro	Arg	Gln
			580					585				590			
Trp	Ala	Arg	His	Asn	Val	Gln	Gly	Thr	Phe	Ser	Asn	Ser	Met	Arg	Tyr

	595					600						605							
Glu	Ala	Phe	Lys	Tyr	Leu	Asp	Ile	Phe	Thr	Ile	Thr	Pro	Glu	Glu	Asn				
	610					615						620							
Asn	Phe	Ala	Phe	Thr	Ile	Asp	Leu	Glu	Ser	Gly	Gly	Asp	Leu	Phe	Ile				
625					630					635					640				
Asp	Lys	Ile	Glu	Phe	Ile	Pro	Val	Ser	Gly	Ser	Ala	Phe	Glu	Tyr	Glu				
				645					650					655					
Gly	Lys	Gln	Asn	Ile	Glu	Lys	Thr	Gln	Lys	Ala	Val	Asn	Asp	Leu	Phe				
			660					665					670						
Ile	Asn																		

<210> 21

<211> 675

<212> PRT

<213> Bacillus thuringiensis

<400> 21

Met	Asn	Pro	Tyr	Gln	Asn	Lys	Ser	Glu	Cys	Glu	Ile	Leu	Asn	Ala	Pro				
1				5				10					15						
Leu	Asn	Asn	Ile	Asn	Met	Pro	Asn	Arg	Tyr	Pro	Phe	Ala	Asn	Asp	Pro				
			20					25					30						
Asn	Ala	Val	Met	Lys	Asn	Gly	Asn	Tyr	Lys	Asp	Trp	Leu	Asn	Glu	Cys				
			35			40						45							
Asp	Gly	Ile	Thr	Pro	Ser	Ile	Phe	Gly	Thr	Leu	Gly	Val	Leu	Ala	Ser				
50						55					60								
Ile	Val	Ile	Ser	Thr	Ile	Asn	Leu	Ala	Thr	Ser	Pro	Ser	Ile	Gly	Asp				
65					70					75					80				
Ala	Phe	Ala	Leu	Val	Ser	Ser	Ile	Gly	Glu	Tyr	Trp	Pro	Glu	Thr	Lys				
				85				90						95					
Thr	Ser	Phe	Pro	Leu	Ser	Val	Ala	Asp	Val	Asn	Arg	Leu	Ile	Arg	Glu				
			100					105					110						
Ala	Leu	Asp	Gln	Asn	Ala	Ile	Asn	Arg	Ala	Thr	Gly	Lys	Phe	Asn	Gly				
			115					120					125						
Leu	Met	Asp	Thr	Tyr	Asn	Thr	Val	Tyr	Leu	Lys	Asn	Leu	Gln	Asp	Trp				
130						135						140							
Tyr	Asp	Thr	Arg	Ile	Pro	Ala	Asn	Pro	Gln	Gly	Asp	Ser	Gln	Leu	Arg				
145					150					155					160				
Glu	Ala	Ala	Arg	Arg	Ser	Leu	Glu	Glu	Ile	Glu	Arg	Asp	Phe	Arg	Lys				
				165						170				175					
Ala	Leu	Ala	Gly	Glu	Phe	Ala	Glu	Ala	Gly	Ser	Gln	Ile	Val	Leu	Leu				
			180					185					190						
Pro	Ile	Tyr	Ala	Gln	Ala	Ala	Asn	Ile	His	Leu	Leu	Ile	Leu	Lys	Asp				
			195					200					205						
Ala	Met	Gln	Phe	Arg	Thr	Asp	Leu	Gly	Leu	Ile	Arg	Pro	Val	Gly	Val				
210						215						220							
Pro	Ile	Thr	Thr	Ser	Ala	Glu	Asp	Pro	Phe	Glu	Ser	Glu	Phe	Leu	Leu				
225					230						235				240				
Arg	Ile	Lys	Lys	Tyr	Thr	Asp	His	Cys	Ile	Ser	Tyr	Tyr	Asp	Asp	Gly				
				245						250				255					
Leu	Ala	Lys	Ile	Arg	Ser	Arg	Gly	Ser	Asp	Gly	Glu	Thr	Trp	Trp	Glu				
			260					265					270						
Phe	Asn	Lys	Phe	Arg	Arg	Glu	Met	Thr	Leu	Thr	Val	Leu	Asp	Leu	Val				
			275					280					285						
Ala	Leu	Tyr	Pro	Thr	His	Asn	Ile	Lys	Leu	Tyr	Pro	Ile	Pro	Thr	Gln				
290						295						300							

Thr Glu Leu Ser Arg Val Val Tyr Thr Asp Pro Val Gly Cys Phe Gly
 305 310 315 320
 Asn Arg Lys Ser Asp Ile Phe Ser Arg Leu Asn Phe Asp Tyr Leu Glu
 325 330 335
 Asn Arg Leu Thr Arg Pro Arg Glu Pro Phe Asn Tyr Leu Asn Ser Val
 340 345 350
 Gln Leu Phe Ala Ser Thr Val Ser Asn Ser Asn Asn Gly Glu Val Leu
 355 360 365
 Arg Gly Asn Leu Asn Lys Ile Met Phe Glu Gly Gly Trp Thr Ala Ser
 370 375 380
 Arg Ser Gly Asp Gly Val Thr Thr Gly Thr Pro Phe Ser Thr Met Asp
 385 390 395 400
 Trp Ser Tyr Gly Trp Gly Tyr Pro Arg Lys His Tyr Ala Glu Ile Thr
 405 410 415
 Ser Arg Ser Gln Ala Leu Pro Gly Leu Asn Asn Ser Ile His Val Ile
 420 425 430
 Val Gly Ile Asp Ser Phe Arg Ala Ile Gly Pro Gly Gly Gln Gly Asp
 435 440 445
 His Thr Phe Ser Leu Pro Gly Gly Asp Met Tyr Asp Cys Gly Lys Val
 450 455 460
 Gln Ile Asn Pro Leu Glu Asp Tyr Arg Asn Ser Asp His Trp Ile Ser
 465 470 475 480
 Asp Met Met Thr Ile Asn Gln Ser Val Gln Leu Ala Ser Asn Pro Thr
 485 490 495
 Gln Thr Phe Ala Phe Ser Ala Leu Ser Leu Gly Trp His His Ser Ser
 500 505 510
 Ala Gly Asn Arg Asn Val Tyr Val Tyr Asp Lys Ile Thr Gln Ile Pro
 515 520 525
 Ala Thr Lys Thr Val Arg Glu His Pro Met Ile Lys Gly Pro Gly Phe
 530 535 540
 Thr Gly Gly Asp Leu Ala Asp Leu Ser Ser Asn Ser Asp Ile Leu Gln
 545 550 555 560
 Tyr Asp Leu Arg Ser Asp Tyr Asp Asp Arg Leu Thr Glu Asp Val Pro
 565 570 575
 Phe Arg Ile Arg Ile Arg Cys Ala Ser Ile Gly Val Ser Thr Ile Ser
 580 585 590
 Val Asp Asn Trp Gly Ser Ser Ser Pro Gln Val Thr Val Ala Ser Thr
 595 600 605
 Ala Ala Ser Leu Asp Thr Leu Lys Tyr Glu Ser Phe Gln Tyr Val Ser
 610 615 620
 Ile Pro Gly Asn Tyr Tyr Phe Asp Ser Ala Pro Arg Ile Arg Leu Leu
 625 630 635 640
 Arg Gln Pro Gly Arg Leu Leu Val Asp Arg Ile Glu Ile Ile Pro Val
 645 650 655
 Asn Phe Phe Pro Leu Ser Glu Gln Glu Asn Lys Ser Val Asp Ser Leu
 660 665 670
 Phe Ile Asn
 675

<210> 22

<211> 659

<212> PRT

<213> *Bacillus thuringiensis*

<400> 22

Asn Ser Tyr Glu Asn Lys Asn Glu Tyr Glu Ile Leu Asn Asp Ser Lys

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Lys	Ser	Asn	Met	Ser	Asn	Pro	Tyr	Leu	Arg	Tyr	Pro	Leu	Ala	Asn	Asp
			20					25					30		
Ser	Leu	Ala	Ser	Met	Gln	Asn	Thr	Asn	Tyr	Lys	Asp	Trp	Leu	Thr	Met
		35					40					45			
Cys	Asp	Arg	Thr	Asp	Thr	Asp	Val	Leu	Ser	Ser	Arg	Gly	Ala	Val	Ser
	50					55				60					
Thr	Gly	Val	Gly	Met	Leu	Ser	Thr	Ile	Leu	Ser	Leu	Phe	Gly	Ile	Pro
65					70					75					80
Leu	Ile	Gly	Glu	Gly	Ile	Asp	Leu	Leu	Leu	Gly	Ala	Ala	Asp	Phe	Leu
				85				90						95	
Trp	Pro	Glu	Ser	Asp	Gly	Gly	His	Gln	Tyr	Thr	Trp	Glu	Asp	Leu	Met
			100					105					110		
Asn	His	Ile	Glu	Glu	Leu	Met	Asp	Glu	Arg	Leu	Glu	Thr	Glu	Lys	Arg
		115					120					125			
Thr	Thr	Ala	Leu	Asp	Asp	Leu	Arg	Gly	Leu	Lys	Ala	Leu	Leu	Gly	Leu
	130					135					140				
Phe	Arg	Asp	Ala	Phe	Asp	Ser	Trp	Glu	Lys	Asn	Gln	Asn	Asp	Pro	Ile
145					150					155					160
Ala	Lys	Asn	Arg	Val	Gly	Gly	Tyr	Phe	Glu	Asp	Val	His	Thr	His	Phe
				165				170						175	
Val	Lys	Asp	Met	Ala	Ser	Ile	Phe	Ser	Ala	Thr	Asn	Tyr	Glu	Val	Leu
			180					185					190		
Leu	Leu	Pro	Val	Tyr	Ala	Gln	Ala	Ala	Asn	Leu	His	Leu	Leu	Leu	Leu
		195					200					205			
Arg	Glu	Gly	Val	Ile	Tyr	Gly	Ser	Arg	Trp	Gly	Ile	Ala	Pro	Ala	Ala
	210					215					220				
Asp	Phe	Tyr	His	Asp	Gln	Leu	Leu	Lys	Tyr	Thr	Ala	Ile	Tyr	Ala	Asn
225					230					235					240
His	Cys	Val	Thr	Trp	Tyr	Asn	Asn	Gly	Leu	Ala	Gln	Gln	Lys	Glu	Leu
				245				250						255	
Phe	Ala	Lys	Ser	Pro	Asn	Trp	Asn	Arg	Phe	Asn	Ala	Tyr	Arg	Arg	Asp
			260					265					270		
Met	Thr	Ile	Thr	Val	Leu	Asp	Ile	Ile	Ala	Leu	Phe	Pro	Thr	Tyr	Asp
		275					280					285			
Ala	Arg	Leu	Tyr	Thr	Lys	Pro	Ile	Lys	Thr	Glu	Leu	Thr	Arg	Glu	Ile
	290					295				300					
Tyr	Ser	Asp	Val	Leu	Asn	Leu	Asp	Val	Tyr	Gly	Val	Gln	Gln	Thr	Asp
305					310					315					320
Leu	Asn	Lys	Asn	Glu	Ala	Ala	Phe	Thr	Arg	Ser	Pro	His	Leu	Val	Thr
				325				330						335	
Arg	Leu	Arg	Gly	Phe	Asp	Phe	Tyr	Thr	Arg	Thr	Lys	Tyr	Ala	Tyr	Trp
			340					345					350		
Arg	Tyr	Leu	Ala	Gly	His	Thr	Asn	Tyr	Phe	Ser	Phe	Thr	Gly	Asn	Gly
		355					360					365			
Thr	Ile	Tyr	Ser	Ser	Ser	Phe	Asn	Asn	Trp	Tyr	Asp	Thr	Asp	Met	Thr
	370					375					380				
Lys	Ser	Thr	Ile	Asn	Ile	Pro	Asp	Tyr	Ala	Asn	Ile	Tyr	Lys	Leu	Trp
385					390					395					400
Thr	Lys	Ser	Tyr	Thr	Asn	Ile	Ser	Pro	Tyr	Thr	Asp	Pro	Val	Gly	Ile
				405				410						415	
Ser	Gln	Met	Gln	Phe	Ser	Leu	Thr	Asn	Asn	Gln	Gln	Leu	Thr	Tyr	Thr
			420					425				430			
Gly	Thr	Ser	Ala	Pro	Lys	Tyr	Pro	Val	Arg	Glu	Thr	Phe	Phe	Glu	Ile
		435					440					445			
Pro	Pro	Thr	Asp	Glu	Lys	Pro	Leu	Thr	Tyr	Glu	Asn	Tyr	Ser	His	Ile
	450					455					460				

Leu	Ser	Tyr	Met	Thr	Ser	Ala	Gln	His	Phe	Gly	Asp	Lys	Lys	Ile	Gly
465					470					475					480
Tyr	Thr	Phe	Ala	Trp	Met	His	Glu	Ser	Val	Asp	Phe	Asp	Asn	Arg	Val
				485					490					495	
Asp	Pro	Asp	Lys	Ile	Thr	Gln	Ile	Pro	Ala	Val	Lys	Gly	Asp	Tyr	Leu
			500					505					510		
Gln	Tyr	Gly	Tyr	Val	Lys	Gln	Gly	Pro	Gly	His	Thr	Gly	Gly	Asp	Leu
		515					520					525			
Val	Ser	Met	Ile	Arg	Thr	Asp	Arg	Leu	Gly	Ile	Asn	Val	Tyr	Phe	Pro
	530					535					540				
Gln	Pro	Leu	Asp	Tyr	Arg	Ile	Arg	Ile	Arg	Tyr	Ser	Thr	Ser	Ser	Asn
545					550					555					560
Gly	Tyr	Leu	Tyr	Ile	Tyr	Ser	Pro	Asn	Thr	Lys	Ile	Val	Tyr	Leu	Pro
				565					570					575	
Pro	Thr	Thr	Leu	Val	Asp	Gly	Gln	Pro	Thr	Phe	Asp	Pro	Met	Asp	Phe
			580					585					590		
Ser	Ala	Phe	Arg	Val	Val	Glu	Val	Pro	Ala	Ser	Phe	Arg	Ala	Ser	Val
		595					600					605			
Ala	Gly	Tyr	Thr	Asn	Phe	Thr	Ile	Glu	Ala	Gly	Phe	Gly	Pro	Val	Tyr
	610					615					620				
Ile	Asp	Lys	Ile	Glu	Phe	Ile	Pro	Asp	Asn	Thr	Thr	Thr	Leu	Glu	Tyr
625					630					635					640
Glu	Gly	Gly	Arg	Asp	Leu	Glu	Lys	Thr	Lys	Asn	Ala	Val	Asn	Asp	Leu
				645					650					655	
Phe	Thr	Asn													

<210> 23

<211> 666

<212> PRT

<213> *Bacillus thuringiensis*

<400> 23

Asn	Ser	Tyr	Glu	Asn	Lys	Asn	Glu	Tyr	Glu	Ile	Leu	Glu	Ser	Ser	Ser
1				5					10					15	
Asn	Asn	Thr	Asn	Met	Pro	Asn	Arg	Tyr	Pro	Phe	Ala	Asn	Asp	Arg	Asp
			20					25				30			
Met	Ser	Thr	Met	Ser	Phe	Asn	Asp	Cys	Gln	Gly	Ile	Ser	Trp	Asp	Glu
		35				40					45				
Ile	Trp	Glu	Ser	Ala	Glu	Thr	Ile	Thr	Ser	Ile	Gly	Ile	Asp	Leu	Ile
	50					55					60				
Glu	Phe	Leu	Met	Glu	Pro	Ser	Leu	Gly	Gly	Ile	Asn	Thr	Leu	Phe	Ser
65				70						75				80	
Ile	Ile	Gly	Lys	Leu	Ile	Pro	Thr	Asn	His	Gln	Ser	Val	Ser	Ala	Leu
			85					90						95	
Ser	Ile	Cys	Asp	Leu	Leu	Ser	Ile	Ile	Arg	Lys	Glu	Val	Ala	Asp	Ser
			100					105					110		
Val	Leu	Ser	Asp	Ala	Ile	Cys	Arg	Phe	Leu	Asp	Gly	Lys	Leu	Lys	Asn
		115					120					125			
Tyr	Arg	Glu	Tyr	Tyr	Leu	Pro	Tyr	Leu	Glu	Ala	Trp	Leu	Lys	Asp	Gly
	130					135					140				
Lys	Pro	Leu	Gln	Lys	Thr	Asn	Asn	Ser	Asp	Ile	Gly	Gln	Leu	Val	Lys
145				150						155					160
Tyr	Phe	Glu	Leu	Ser	Glu	Arg	Asp	Phe	Asn	Glu	Ile	Leu	Gly	Gly	Ser
				165					170					175	
Leu	Ala	Arg	Asn	Asn	Ala	Gln	Ile	Leu	Leu	Leu	Pro	Tyr	Phe	Cys	Ala

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Asn	Ser	Thr	Ala	Leu	Glu	Tyr	Glu	Gly	Lys	Gln	Ser	Leu	Glu	Lys	Ala
				645					650					655	
Gln	Asp	Val	Val	Asn	Asp	Leu	Phe	Val	Lys						
			660					665							